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PCT/EP00/02410

SEQUENZPROTOKOLL

<110> Biosyn Arzneimittel GmbH

<120> Nukleinsäuremolekül, umfassend eine für ein Hämocyanin kodierende Nukleinsäuresequenz

<130> PCT1153-01966

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<170> PatentIn Ver. 2.1

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4

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<213> *Haliotis tuberculata*

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5

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<213> *Haliotis tuberculata*

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<213> *Haliotis tuberculata*

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 gattttgcg gcatggatt cagcagtctg cactagttaa attcttgcg tgcaaatc 1020
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 catggggcata tgatcgctt taaaatatg agatcactga gcagctcaat gcccggatc 1140
 tacacatcgg agatagattt ttcatcagat acgaagcgtt tgatcttcat ggtacaagtc 1200
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<210> 11

<211> 1244

<212> DNA

<213> *Haliotis tuberculata*

<400> 11

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<210> 12

<211> 1255

<212> DNA

<213> *Haliotis tuberculata*

<400> 12

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 ccatctgcac ctcaccgtt tgctgtgtt gtccacggca tggctacatt tccccagtt 240
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 cacccatcgat atgatgatga cttcacaatc tctgtcaatc tgaccgccaa caacggact 1200
 gtcctgagca gcagtctaat cccaaacacccg agtgtcatat tccagcgggg acatc 1255

<210> 13

<211> 1248

<212> DNA

<213> *Haliotis tuberculata*

<400> 13

gtgacataaa taccaggaggc atgtcaccga accgtgttcg ccgtgagctg agcgatctgt 60
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 ccaacggata ccaggcttcc gcagccttcc atggcttacc agcaggctgc catgatagcc 180
 gggaaatga gatcgcatgt tgcatcaccg ggatgcccac cttccccccag tggcacagac 240
 tgtacaccct gcagttggag atgctctga ggagacatgg atcatctgtc gccatccct 300
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 caaataccta cacagtaaga gaccacagg agatgtgtt ccagtttgcgaaatggag 480
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 tgaatcagaa cccattcaca aagatcaccg cagttccaa cacatctat gactacgaga 840
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<210> 14

<211> 1207

<212> DNA

<213> *Haliotis tuberculata*

<400> 14

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 atgatactgg tcccaatgtt taccatggca tagcatctt ccacggaaatgttccatgt 180
 gcgagatgaa cggccgcacccat gttgcctgtt gtgtcaccgg tatggcctcc ttcccacact 240
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 cccgagaacaa tcccttcat gagggtcgatc ttgtatcatc cgggtgtacc acgtcagtt 420
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 aagtcccttgc ggctttggag cagactgact actgcccattt cgaagtccatc tttgagatgt 540
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 aacttgg 1207

<210> 15

<211> 1546

<212> DNA

<213> *Haliotis tuberculata*

<400> 15

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 acactttcat cgtaaggaag ttgattcgct gtccctggat gaagcaaaca acttgaagaa 120
 tgcccttac aagctacaga acgaccacag tctaacggga tacgaagcaa tctctggta 180
 ccatggatac cccaatctgt gtccggaaaga aggcatgac aaaaatcccc tgctgcgtcc 240
 ccggatgggc atcttcctt actggcacag actcttgcacc attcaactgg aaagagctct 300
 tgagcacaat ggtgcactgc ttgtgttcc ttactggac tggaaacaagg acctgtcgctc 360
 actgcccggcg ttcttctccg actccagcaa caacaatccc tacttcaagt accacatcgc 420
 cggtgttggt cacgacacccg tcaagagagcc aactagtctt atatataacc agccccaaat 480
 ccatggttat gattatctt attacatgc attgaccacg cttaaagaaa acaattactg 540
 ggacttttag gttcagtagt agatccctca caacggcgtc cactctggc ttggaggatc 600
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 ccacgtcttat tctgtcaagc ctgtgtacta ctatgttact ggacccacga gagacctttg 1500
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<210> 16

<211> 967

<212> DNA

<213> *Megathura crenulata*

<400> 16

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 aaaacttatg tggatttcac tggatgcatttcc cacacaatc cttttcatag ttcatgttatt 120
 gcatttgaag aaaatgtcc ccacaccaaa agacaaatag atcaaagact ctttaaaaccc 180
 gctaccttgc gacaccacac agacctgttc aaccagattt tgatgcctt tgaacaagaa 240
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 tactttcacc attctaacttgc ttatgttctt tggttgcgtt ggcaagcctt acagatgaga 420

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 aagatctacg actatgaaaaa tgcctccat tacacatacg aagatccaac atttgaggc 600
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 aagtggggat ttgatcgcgt ttcaagttt gacatcacgc acgtttgaa agatctcgat 840
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 cttgcattca gtcttattcc acatgcttct gtcattcgtg agcatgcacg tggtaagctg 960
 aatagag 967

<210> 17
 <211> 1242
 <212> DNA
 <213> Megathura crenulata

<400> 17
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 ccgaggagat gaatgaactt cgtaaagccc tagcctact gaaagaggac aaaagtgcgg 120
 gtggatttca gcagcttggt gcattccatg gggagccaaa atgggtgcctt agtcccgaag 180
 cacttaaaaa atttgcctgc tttgttcacg gcatgtctgt gtcccttcac tggcatcgac 240
 tttgtgacggt tcagagtgaa aatgttttga gacgacatgg ctacgatgg gctttgcgt 300
 actgggattt gaccttcctt cttaatcacc ttcccaactt ggcagatcat gagaagtacg 360
 tcgaccctga agatggggta gagaagcata acccttgggtt cgatggtcat atagatacag 420
 tcgacaaaac aacaacaaga agtggtcaga ataaacttctt cgaacagcctt gagttggc 480
 attatacaga cattgcggaa caagtactgc tagcgttgg aacaggacaat ttctgtact 540
 ttgaaatcca atatgagatt gcccataact acatccatgc acttggtagga ggcgctcagc 600
 cttatggat ggcatcgctt cgctacactg ctttgcattt actattcttac ttgcattact 660
 ctaatacaga tcgttatgg gcaatatggc aggctttaca gaagtacaga ggaaaaccgt 720
 acaacgttgc taactgttgc ttacatcga tgagagaaacc ttgcacacca ttggccctct 780
 ctgccaatcat caacacacag catgttacca aggagcatcc agtgcattt aacgttttg 840
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<210> 18
 <211> 1236
 <212> DNA
 <213> Megathura crenulata

<400> 18
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 acattgaaaaa tctgaggcctt ggtgaactcg aaagtctgg agtgccttc ctggaaattt 120
 aaaaacgatgg aacttacgaa tcaatagctt aattccatgg tagccctgg tttgtccagt 180
 taaatggtaa ccccatctt tttgtgttcc atggcatgccc aactttccctt cactggcaca 240
 gactgtacgt gttgtcggtt gagaatggccc ttctggaaaaa agatcatctt gtagctttc 300
 cctattggga ctggacaaaaa cgaatcgaaat atttacatca cctgatttca gacgcccattt 360
 actacaatttcc caggcaacat cactatgaga caaaaccattt ccatcatggc aaaaatcacac 420
 acgagaatgaa aatctacttactt agggatccca aggacagcctt ctccatcttca gactactttt 480
 acgagcagggt ctttacgccc ttggagcaggataaacttctt tgatttgcgg attcagttgg 540
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10

gcgctatcca attgatgcac acgccactcc agccgttga taagagcgac aacaatgacg 780
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 tccactctcc cactatactg tttgaggccg gaacag 1236

<210> 19

<211> 241

<212> DNA

<213> Megathura crenulata

<400> 19

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 aattggacaa gctcaacaa ctgtcaactgg tgaaagccct cgagtccatg aaagccgacc 120
 attcatctga tgggttccag gcaatcgctt ccttcatgc tcttcctt ctttgtccat 180
 caccagctgc ttcaaaaggagg tttgcgtgt gcgtccatgg catgccaacc ttcccgaat 240
 g 241

<210> 20

<211> 949

<212> DNA

<213> Megathura crenulata

<400> 20

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 gagacctacc tcgatccatg tactgggaa actaaaaaca acccttcca tcacgccc 120
 gtggcgtttggaaatgggtt aacaaggcagg aatcctgtatg cccaaactttt tatgaaacca 180
 acttacggag accacactta ccttcgtcacc agcatgtatc acgcatttgc gcagggaaagac 240
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 ggcaagtctt acaaggccca ctgcgcctcg tctcaagaaa gagaaccatt aaagcctttt 480
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<210> 21

<211> 760

<212> DNA

<213> Megathura crenulata

<400> 21

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11

actgggactg gactcgatca atgagcgccc ttccacattt tggctgtat cctacttaca 360
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 ttggcatga tactacaaga gatgtcggt atgatctta tcaatctcct ggttccgtc 480
 actacacaga tattgcacaa caagtcctc tggccttga gcaggacagt ttctgtgatt 540
 ttgaggtaca atttgaatt gcccataatt tcatacatgc actgatttgtt ggtaacgaac 600
 catacagtt gtcatcttg aggtatacta catacgatcc aatcttctt ttgcaccact 660
 ccagtaga ccgactttgg gccatctggc aagcaatcac tagtgcggcc gcctgcaggt 720
 cgaccataag ggagagctcc caacgcgttg gatgcaatct 760

<210> 22

<211> 323

<212> DNA

<213> *Megathura crenulata*

<400> 22

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 atgatggaca tcctgttgc tttgtgtcc atggcatgcc cacccttccc cactggcaca 240
 gactgtacgt tcttcagggtt gagaatgcgc tcttagaacg agggctgca gttgctgttc 300
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<210> 23

<211> 988

<212> DNA

<213> *Megathura crenulata*

<400> 23

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 cccgccttag tgactgacca cgagaacaat cccttccacc acggccatat tggctcatctg 180
 aatgtggata catctcgatc tccaaagagac atgctgttta atgatcctga acaaggctca 240
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 gaagttcagt ttgaacttac acacaatgcc atccactt ggactggagg acataactcca 360
 tatgaaatgt catcaacttggaa atatacagca tatgatccac tcttttatct ccaccattcc 420
 aacactgatc gtatctggc catctggcag gcactccaga aatatacgagg tcttccatata 480
 aacgcagctc actgcgatata ccaagttctg aaacaaccctt taaaaccatt cagcgagttcc 540
 aggaatccaa acccagtca cagaqccaaat tctaggcccg ttgatttatt tgattatgag 600
 aaattcaattt atcaatatga cacacttacc ttccacggac tttctatccc agaacttgat 660
 gccatgcttc aagagagaaa gaaggaagag agaacatttgc cagcccttctt gttgcacgg 720
 tttggcgcctt gtgctgtatgt ttcgtttgtat gtctgcacac ctgtatggta ttgtgccttt 780
 gctggAACCT tcgcgttact tgggtggggag cttgagatgc cctggccctt tggaaatgg 840
 ttccgttacg atatcacaaa gtttctcaag cagatgaatc ttcaactatga ttctgagttc 900
 cactttgagt tgaagattgt tggcacagat ggaacagaac tgccatcgaa tcgtatcaag 960
 agcccttacca ttgaacacca tggaggag 988

<210> 24

<211> 310

<212> DNA

<213> *Megathura crenulata*

<400> 24

gtcacgatca cagtgaacgt cacgatggat ttttcaggaa ggaagtcggg tccctgtccc 60
 tggatgaagc caatgacccctt aaaaatgcac ttttacaatgtt gcagaatgtt cagggtccca 120
 atggatatga atcaatagcc gtttaccatg gtttaccatc ccttgcctt gaacatgggt 180
 aagaccagta cgcacgtgtt gtttacccatg ttttacccatc ctttgcctt ctttgcctt 240
 atacaatcca gtttggatgtt gtttacccatg aacatgggtt tcattttgggtt ctggccatact 300

12

310

gggactggac

<210> 25

<211> 422

<212> PRT

<213> Haliotis tuberculata

<220>

<221> SIGNAL

<222> (1)..(15)

<400> 25

Leu Val Gln Phe Leu Leu Val Ala Leu Val Ala Gly Ala Gly Ala Asp
1 5 10 15Asn Val Val Arg Lys Asp Val Ser His Leu Thr Asp Asp Glu Val Gln
20 25 30Ala Leu His Gly Ala Leu His Asp Val Thr Ala Ser Thr Gly Pro Leu
35 40 45Ser Phe Glu Asp Ile Thr Ser Tyr His Ala Ala Pro Ala Ser Cys Asp
50 55 60Tyr Lys Gly Arg Lys Ile Ala Cys Cys Val His Gly Met Pro Ser Phe
65 70 75 80Pro Phe Trp His Arg Ala Tyr Val Val Gln Ala Glu Arg Ala Leu Leu
85 90 95Ser Lys Arg Lys Thr Val Gly Met Pro Tyr Trp Asp Trp Thr Gln Thr
100 105 110Leu Thr His Leu Pro Ser Leu Val Thr Glu Pro Ile Tyr Ile Asp Ser
115 120 125Lys Gly Gly Lys Ala Gln Thr Asn Tyr Trp Tyr Arg Gly Glu Ile Ala
130 135 140Phe Ile Asn Lys Lys Thr Ala Arg Ala Val Asp Asp Arg Leu Phe Glu
145 150 155 160Lys Val Glu Pro Gly His Tyr Thr His Leu Met Glu Thr Val Leu Asp
165 170 175Ala Leu Glu Gln Asp Glu Phe Cys Lys Phe Glu Ile Gln Phe Glu Leu
180 185 190Ala His Asn Ala Ile His Tyr Leu Val Gly Gly Lys Phe Glu Tyr Ser
195 200 205Met Ser Asn Leu Glu Tyr Thr Ser Tyr Asp Pro Ile Phe Phe Leu His
210 215 220His Ser Asn Val Asp Arg Leu Phe Ala Ile Trp Gln Arg Leu Gln Glu
225 230 235 240

13

Leu Arg Gly Lys Asn Pro Asn Ala Met Asp Cys Ala His Glu Leu Ala
 245 250 255

His Gln Gln Leu Gln Pro Phe Asn Arg Asp Ser Asn Pro Val Gln Leu
 260 265 270

Thr Lys Asp His Ser Thr Pro Ala Asp Leu Phe Asp Tyr Lys Gln Leu
 275 280 285

Gly Tyr Ser Tyr Asp Ser Leu Asn Leu Asn Gly Met Thr Pro Glu Gln
 290 295 300

Leu Lys Thr Glu Leu Asp Glu Arg His Ser Lys Glu Arg Ala Phe Ala
 305 310 315 320

Ser Phe Arg Leu Ser Gly Phe Gly Ser Ala Asn Val Val Val Tyr
 325 330 335

Ala Cys Val Pro Asp Asp Asp Pro Arg Ser Asp Asp Tyr Cys Glu Lys
 340 345 350

Ala Gly Asp Phe Phe Ile Leu Gly Gly Gln Ser Glu Met Pro Trp Arg
 355 360 365

Phe Tyr Arg Pro Phe Phe Tyr Asp Val Thr Glu Ala Val His His Leu
 370 375 380

Gly Val Pro Leu Ser Gly His Tyr Tyr Val Lys Thr Glu Leu Phe Ser
 385 390 395 400

Val Asn Gly Thr Ala Leu Ser Pro Asp Leu Leu Pro Gln Pro Thr Val
 405 410 415

Ala Tyr Arg Pro Gly Lys
 420

<210> 26

<211> 419

<212> PRT

<213> Haliotis tuberculata

<400> 26

Gly His Leu Asp Pro Pro Val His His Arg His Asp Asp Asp Leu Ile
 1 5 10 15

Val Arg Lys Asn Ile Asp His Leu Thr Arg Glu Glu Glu Tyr Glu Leu
 20 25 30

Arg Met Ala Leu Glu Arg Phe Gln Ala Asp Thr Ser Val Asp Gly Tyr
 35 40 45

Gln Ala Thr Val Glu Tyr His Gly Leu Pro Ala Arg Cys Pro Arg Pro
 50 55 60

Asp Ala Lys Val Arg Phe Ala Cys Cys Met His Gly Met Ala Ser Phe
 65 70 75 80

Pro His Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Val
 85 90 95

Arg Arg Gly Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Lys Pro
 100 105 110

Met Thr His Leu Pro Asp Leu Ala Ser Asn Glu Thr Tyr Val Asp Pro
 115 120 125

Tyr Gly His Thr His His Asn Pro Phe Phe Asn Ala Asn Ile Ser Phe
 130 135 140

Glu Glu Gly His His His Thr Ser Arg Met Ile Asp Ser Lys Leu Phe
 145 150 155 160

Ala Pro Val Ala Phe Gly Glu His Ser His Leu Phe Asp Gly Ile Leu
 165 170 175

Tyr Ala Phe Glu Gln Glu Asp Phe Cys Asp Phe Glu Ile Gln Phe Glu
 180 185 190

Leu Val His Asn Ser Ile His Ala Trp Ile Gly Gly Ser Glu Asp Tyr
 195 200 205

Ser Met Ala Thr Leu His Tyr Thr Ala Phe Asp Pro Ile Phe Tyr Leu
 210 215 220

His His Ser Asn Val Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln
 225 230 235 240

Ile Arg Arg His Lys Pro Tyr Gln Ala His Cys Ala Gln Ser Val Glu
 245 250 255

Gln Leu Pro Met Lys Pro Phe Ala Phe Pro Ser Pro Leu Asn Asn Asn
 260 265 270

Glu Lys Thr His Ser His Ser Val Pro Thr Asp Ile Tyr Asp Tyr Glu
 275 280 285

Glu Val Leu His Tyr Ser Tyr Asp Asp Leu Thr Phe Gly Gly Met Asn
 290 295 300

Leu Glu Glu Ile Glu Glu Ala Ile His Leu Arg Gln Gln His Glu Arg
 305 310 315 320

Val Phe Ala Gly Phe Leu Leu Ala Gly Ile Gly Thr Ser Ala Leu Val
 325 330 335

Asp Ile Phe Ile Asn Lys Pro Gly Asn Gln Pro Leu Lys Ala Gly Asp
 340 345 350

Ile Ala Ile Leu Gly Gly Ala Lys Glu Met Pro Trp Ala Phe Asp Arg
 355 360 365

Leu Tyr Lys Val Glu Ile Thr Asp Ser Leu Lys Thr Leu Ser Leu Asp
 370 375 380

15

Val Asp Gly Asp Tyr Glu Val Thr Phe Lys Ile His Asp Met His Gly
385 390 395 400

Asn Ala Leu Asp Thr Asp Leu Ile Pro His Ala Ala Val Val Ser Glu
405 410 415

Pro Ala His

<210> 27

<211> 414

<212> PRT

<213> Haliotis tuberculata

<400> 27

Pro Thr Phe Glu Asp Glu Lys His Ser Leu Arg Ile Arg Lys Asn Val
1 5 10 15

Asp Ser Leu Thr Pro Glu Glu Thr Asn Glu Leu Arg Lys Ala Leu Glu
20 25 30

Leu Leu Glu Asn Asp His Thr Ala Gly Gly Phe Asn Gln Leu Gly Ala
35 40 45

Phe His Gly Glu Pro Lys Trp Cys Pro Asn Pro Glu Ala Glu His Lys
50 55 60

Val Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
65 70 75 80

Leu Leu Ala Leu Gln Ala Glu Asn Ala Leu Arg Lys His Gly Tyr Ser
85 90 95

Gly Ala Leu Pro Tyr Trp Asp Trp Thr Arg Pro Leu Ser Gln Leu Pro
100 105 110

Asp Leu Val Ser His Glu Gln Tyr Thr Asp Pro Ser Asp His His Val
115 120 125

Lys His Asn Pro Trp Phe Asn Gly His Ile Asp Thr Val Asn Gln Asp
130 135 140

Thr Thr Arg Ser Val Arg Glu Asp Leu Tyr Gln Gln Pro Glu Phe Gly
145 150 155 160

His Phe Thr Asp Ile Ala Gln Gln Val Leu Leu Ala Leu Glu Gln Asp
165 170 175

Asp Phe Cys Ser Phe Glu Val Gln Tyr Glu Ile Ser His Asn Phe Ile
180 185 190

His Ala Leu Val Gly Gly Thr Asp Ala Tyr Gly Met Ala Ser Leu Arg
195 200 205

16

Tyr Thr Ala Tyr Asp Pro Ile Phe Phe Leu His His Ser Asn Thr Asp
 210 215 220
 Arg Ile Trp Ala Ile Trp Gln Ser Leu Gln Lys Tyr Arg Gly Lys Pro
 225 230 235 240
 Tyr Asn Thr Ala Asn Cys Ala Ile Glu Ser Met Arg Arg Pro Leu Gln
 245 250 255
 Pro Phe Gly Leu Ser Ser Ala Ile Asn Pro Asp Arg Ile Thr Arg Glu
 260 265 270
 His Ala Ile Pro Phe Asp Val Phe Asn Tyr Arg Asp Asn Leu His Tyr
 275 280 285
 Val Tyr Asp Thr Leu Glu Phe Asn Gly Leu Ser Ile Ser Gln Leu Asp
 290 295 300
 Arg Glu Leu Glu Lys Ile Lys Ser His Glu Arg Val Phe Ala Gly Phe
 305 310 315 320
 Leu Leu Ser Gly Ile Lys Lys Ser Ala Leu Val Lys Phe Glu Val Cys
 325 330 335
 Thr Pro Pro Asp Asn Cys His Lys Ala Gly Glu Phe Tyr Leu Leu Gly
 340 345 350
 Asp Glu Asn Glu Met Ala Trp Ala Tyr Asp Arg Leu Phe Lys Tyr Asp
 355 360 365
 Ile Thr Gln Val Leu Glu Ala Asn His Leu His Phe Tyr Asp His Leu
 370 375 380
 Phe Ile Arg Tyr Glu Val Phe Asp Leu Lys Gly Val Ser Leu Gly Thr
 385 390 395 400
 Asp Leu Phe His Thr Ala Asn Val Val His Asp Ser Gly Thr
 405 410

<210> 28
 <211> 413
 <212> PRT
 <213> Haliotis tuberculata

<400> 28
 Gly Thr Arg Asp Arg Asp Asn Tyr Val Glu Glu Val Thr Gly Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Leu Asn Asp Leu Asn Thr Gly Glu Met Glu Ser
 20 25 30
 Leu Arg Ala Ala Phe Leu His Ile Gln Asp Asp Gly Thr Tyr Glu Ser
 35 40 45
 Ile Ala Gln Tyr His Gly Lys Pro Gly Lys Cys Gln Leu Asn Asp His
 50 55 60

Asn Ile Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
 65 70 75 80

Arg Leu Tyr Val Val Gln Val Glu Asn Ala Leu Leu Asn Arg Gly Ser
 85 90 95

Gly Val Ala Val Pro Tyr Trp Glu Trp Thr Ala Pro Ile Asp His Leu
 100 105 110

Pro His Phe Ile Asp Asp Ala Thr Tyr Phe Asn Ser Arg Gln Gln Arg
 115 120 125

Tyr Asp Pro Asn Pro Phe Phe Arg Gly Lys Val Thr Phe Glu Asn Ala
 130 135 140

Val Thr Thr Arg Asp Pro Gln Ala Gly Leu Phe Asn Ser Asp Tyr Met
 145 150 155 160

Tyr Glu Asn Val Leu Leu Ala Leu Glu Gln Glu Asn Tyr Cys Asp Phe
 165 170 175

Glu Ile Gln Phe Glu Leu Val His Asn Ala Leu His Ser Met Leu Gly
 180 185 190

Gly Lys Gly Gln Tyr Ser Met Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205

Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220

Trp Gln Glu Leu Gln Arg Phe Arg Glu Leu Pro Tyr Glu Glu Ala Asn
 225 230 235 240

Cys Ala Ile Asn Leu Met His Gln Pro Leu Lys Pro Phe Ser Asp Pro
 245 250 255

His Glu Asn His Asp Asn Val Thr Leu Lys Tyr Ser Lys Pro Gln Asp
 260 265 270

Gly Phe Asp Tyr Gln Asn His Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285

Phe His His Leu Ser Ile Pro Ser Leu Asp Ala Thr Leu Lys Gln Arg
 290 295 300

Arg Asn His Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320

Thr Ser Ala Asp Ile Thr Ile Tyr Ile Cys Leu Pro Asp Gly Arg Arg
 325 330 335

Gly Asn Asp Cys Ser His Glu Ala Gly Thr Phe Tyr Ile Leu Gly Gly
 340 345 350

Glu Thr Glu Met Pro Phe Ile Phe Asp Arg Leu Tyr Lys Phe Glu Ile
 355 360 365

Thr Lys Pro Leu Gln Gln Leu Gly Val Lys Leu His Gly Gly Val Phe
 370 375 380
 Glu Leu Glu Leu Glu Ile Lys Ala Tyr Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400
 His Thr Phe Asp Pro Thr Ile Ile Phe Glu Pro Gly Thr
 405 410

<210> 29
 <211> 420
 <212> PRT
 <213> Haliotis tuberculata

<400> 29
 Asp Thr His Ile Leu Asp His Asp His Glu Glu Glu Ile Leu Val Arg
 1 5 10 15

Lys Asn Ile Ile Asp Leu Ser Pro Arg Glu Arg Val Ser Leu Val Lys
 20 25 30

Ala Leu Gln Arg Met Lys Asn Asp Arg Ser Ala Asp Gly Tyr Gln Ala
 35 40 45

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Asn Pro Ser Ala
 50 55 60

Ala His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln
 65 70 75 80

Trp His Arg Leu Tyr Thr Val Gln Val Gln Asp Ala Leu Arg Arg His
 85 90 95

Gly Ser Leu Val Gly Ile Pro Tyr Trp Asp Trp Thr Lys Pro Val Asn
 100 105 110

Glu Leu Pro Glu Leu Leu Ser Ser Ala Thr Phe Tyr His Pro Ile Arg
 115 120 125

Asn Ile Asn Ile Ser Asn Pro Phe Leu Gly Ala Asp Ile Glu Phe Glu
 130 135 140

Gly Pro Gly Val His Thr Glu Arg His Ile Asn Thr Glu Arg Leu Phe
 145 150 155 160

His Ser Gly Asp His Asp Gly Tyr His Asn Trp Phe Phe Glu Thr Val
 165 170 175

Leu Phe Ala Leu Glu Gln Glu Asp Tyr Cys Asp Phe Glu Ile Gln Phe
 180 185 190

Glu Ile Ala His Asn Gly Ile His Thr Trp Ile Gly Gly Ser Ala Val
 195 200 205

19

Tyr Gly Met Gly His Leu His Tyr Ala Ser Tyr Asp Pro Ile Phe Tyr
 210 215 220

Ile His His Ser Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu
 225 230 235 240

Gln Lys Tyr Arg Gly Leu Ser Gly Ser Glu Ala Asn Cys Ala Ile Glu
 245 250 255

His Met Arg Thr Pro Leu Lys Pro Phe Ser Phe Gly Pro Pro Tyr Asn
 260 265 270

Leu Asn Ser His Thr Gln Glu Tyr Ser Lys Pro Glu Asp Thr Phe Asp
 275 280 285

Tyr Lys Lys Phe Gly Tyr Arg Tyr Asp Ser Leu Glu Leu Glu Gly Arg
 290 295 300

Ser Ile Ser Arg Ile Asp Glu Leu Ile Gln Gln Arg Gln Glu Lys Asp
 305 310 315 320

Arg Thr Phe Ala Gly Phe Leu Leu Lys Gly Phe Gly Thr Ser Ala Ser
 325 330 335

Val Ser Leu Gln Val Cys Arg Val Asp His Thr Cys Lys Asp Ala Gly
 340 345 350

Tyr Phe Thr Ile Leu Gly Gly Ser Ala Glu Met Pro Trp Ala Phe Asp
 355 360 365

Arg Leu Tyr Lys Tyr Asp Ile Thr Lys Thr Leu His Asp Met Asn Leu
 370 375 380

Arg His Glu Asp Thr Phe Ser Ile Asp Val Thr Ile Thr Ser Tyr Asn
 385 390 395 400

Gly Thr Val Leu Ser Gly Asp Leu Ile Gln Thr Pro Ser Ile Ile Phe
 405 410 415

Val Pro Gly Arg
 420

<210> 30
 <211> 417
 <212> PRT
 <213> Haliotis tuberculata

<400> 30
 His Lys Leu Asn Ser Arg Lys His Thr Pro Asn Arg Val Arg His Glu
 1 5 10 15

Leu Ser Ser Leu Ser Ser Arg Asp Ile Ala Ser Leu Lys Ala Ala Leu
 20 25 30

Thr Ser Leu Gln His Asp Asn Gly Thr Asp Gly Tyr Gln Ala Ile Ala
 35 40 45

Ala Phe His Gly Val Pro Ala Gln Cys His Glu Pro Ser Gly Arg Glu
 50 55 60

Ile Ala Cys Cys Ile His Gly Met Ala Thr Phe Pro His Trp His Arg
 65 70 75 80

Leu Tyr Thr Leu Gln Leu Glu Gln Ala Leu Arg Arg His Gly Ser Ser
 85 90 95

Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Pro Ile Thr Glu Leu Pro
 100 105 110

His Ile Leu Thr Asp Gly Glu Tyr Tyr Asp Val Trp Gln Asn Ala Val
 115 120 125

Leu Ala Asn Pro Phe Ala Arg Gly Tyr Val Lys Ile Lys Asp Ala Phe
 130 135 140

Thr Val Arg Asn Val Gln Glu Ser Leu Phe Lys Met Ser Ser Phe Gly
 145 150 155 160

Lys His Ser Leu Leu Phe Asp Gln Ala Leu Leu Ala Leu Glu Gln Thr
 165 170 175

Asp Tyr Cys Asp Phe Glu Val Gln Phe Glu Val Met His Asn Thr Ile
 180 185 190

His Tyr Leu Val Gly Gly Arg Gln Thr Tyr Ala Phe Ser Ser Leu Glu
 195 200 205

Tyr Ser Ser Tyr Asp Pro Ile Phe Phe Ile His His Ser Phe Val Asp
 210 215 220

Lys Ile Trp Ala Val Trp Gln Glu Leu Gln Ser Arg Arg His Leu Gln
 225 230 235 240

Phe Arg Thr Ala Asp Cys Ala Val Gly Leu Met Gly Gln Ala Met Arg
 245 250 255

Pro Phe Asn Lys Asp Phe Asn His Asn Ser Phe Thr Lys Lys His Ala
 260 265 270

Val Pro Asn Thr Val Phe Asp Tyr Glu Asp Leu Gly Tyr Asn Tyr Asp
 275 280 285

Asn Leu Glu Ile Ser Gly Leu Asn Leu Asn Glu Ile Glu Ala Leu Ile
 290 295 300

Ala Lys Arg Lys Ser His Ala Arg Val Phe Ala Gly Phe Leu Leu Phe
 305 310 315 320

Gly Leu Gly Thr Ser Ala Asp Ile His Leu Glu Ile Cys Lys Thr Ser
 325 330 335

Glu Asn Cys His Asp Ala Gly Val Ile Phe Ile Leu Gly Gly Ser Ala
 340 345 350

Glu Met His Trp Ala Tyr Asn Arg Leu Tyr Lys Tyr Asp Ile Thr Glu
 355 360 365

Ala Leu Gln Glu Phe Asp Ile Asn Pro Glu Asp Val Phe His Ala Asp
 370 375 380

Glu Pro Phe Phe Leu Arg Leu Ser Val Val Ala Val Asn Gly Thr Val
 385 390 395 400

Ile Pro Ser Ser His Leu His Gln Pro Thr Ile Ile Tyr Glu Pro Gly
 405 410 415

Glu

<210> 31

<211> 403

<212> PRT

<213> Haliotis tuberculata

<400> 31

Asp His His Asp Asp His Gln Ser Gly Ser Ile Ala Gly Ser Gly Val
 1 5 10 15

Arg Lys Asp Val Asn Thr Leu Thr Lys Ala Glu Thr Asp Asn Leu Arg
 20 25 30

Glu Ala Leu Trp Gly Val Met Ala Asp His Gly Pro Asn Gly Phe Gln
 35 40 45

Ala Ile Ala Ala Phe His Gly Lys Pro Ala Leu Cys Pro Met Pro Asp
 50 55 60

Gly His Asn Tyr Ser Cys Cys Thr His Gly Met Ala Thr Phe Pro His
 65 70 75 80

Trp His Arg Leu Tyr Thr Lys Gln Met Glu Asp Ala Met Arg Ala His
 85 90 95

Gly Ser His Val Gly Leu Pro Tyr Trp Asp Trp Thr Ala Ala Phe Thr
 100 105 110

His Leu Pro Thr Leu Val Thr Asp Thr Asp Asn Asn Pro Phe Gln His
 115 120 125

Gly His Ile Asp Tyr Leu Asn Val Ser Thr Thr Arg Ser Pro Arg Asp
 130 135 140

Met Leu Phe Asn Asp Pro Glu His Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160

Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Phe Cys Lys Phe Glu Val
 165 170 175

22

Gln Phe Glu Ile Thr His Asn Ala Ile His Ser Trp Thr Gly Gly His
 180 185 190

Ser Pro Tyr Gly Met Ser Thr Leu Asp Phe Thr Ala Tyr Asp Pro Leu
 195 200 205

Phe Trp Leu His His Ser Asn Thr Asp Arg Ile Trp Ala Val Trp Gln
 210 215 220

Ala Leu Gln Glu Tyr Arg Gly Leu Pro Tyr Asn His Ala Asn Cys Glu
 225 230 235 240

Ile Gln Ala Met Lys Thr Pro Leu Arg Pro Phe Ser Asp Asp Ile Asn
 245 250 255

His Asn Pro Val Thr Lys Ala Asn Ala Lys Pro Leu Asp Val Phe Glu
 260 265 270

Tyr Asn Arg Leu Ser Phe Gln Tyr Asp Asn Leu Ile Phe His Gly Tyr
 275 280 285

Ser Ile Pro Glu Leu Asp Arg Val Leu Glu Glu Arg Lys Glu Glu Asp
 290 295 300

Arg Ile Phe Ala Ala Phe Leu Leu Ser Gly Ile Lys Arg Ser Ala Asp
 305 310 315 320

Val Val Phe Asp Ile Cys Gln Pro Glu His Glu Cys Val Phe Ala Gly
 325 330 335

Thr Phe Ala Ile Leu Gly Gly Glu Leu Glu Met Pro Trp Ser Phe Asp
 340 345 350

Arg Leu Phe Arg Tyr Asp Ile Thr Lys Val Met Lys Gln Leu His Leu
 355 360 365

Arg His Asp Ser Asp Phe Thr Phe Arg Val Lys Ile Val Gly Thr Asp
 370 375 380

Asp His Glu Leu Pro Ser Asp Ser Val Lys Ala Pro Thr Ile Glu Phe
 385 390 395 400

Glu Pro Gly

<210> 32

<211> 511

<212> PRT

<213> Haliotis tuberculata

<400> 32

Val His Arg Gly Gly Asn His Glu Asp Glu His His Asp Asp Arg Leu
 1 5 10 15

Ala Asp Val Leu Ile Arg Lys Glu Val Asp Phe Leu Ser Leu Gln Glu
 20 25 30

Ala Asn Ala Ile Lys Asp Ala Leu Tyr Lys Leu Gln Asn Asp Asp Ser
 35 40 45

Lys Gly Gly Phe Glu Ala Ile Ala Gly Tyr His Gly Tyr Pro Asn Met
 50 55 60

Cys Pro Glu Arg Gly Thr Asp Lys Tyr Pro Cys Cys Val His Gly Met
 65 70 75 80

Pro Val Phe Pro His Trp His Arg Leu His Thr Ile Gln Met Glu Arg
 85 90 95

Ala Leu Lys Asn His Gly Ser Pro Met Gly Ile Pro Tyr Trp Asp Trp
 100 105 110

Thr Lys Lys Met Ser Ser Leu Pro Ser Phe Phe Gly Asp Ser Ser Asn
 115 120 125

Asn Asn Pro Phe Tyr Lys Tyr Tyr Ile Arg Gly Val Gln His Glu Thr
 130 135 140

Thr Arg Asp Val Asn Gln Arg Leu Phe Asn Gln Thr Lys Phe Gly Glu
 145 150 155 160

Phe Asp Tyr Leu Tyr Tyr Leu Thr Leu Gln Val Leu Glu Glu Asn Ser
 165 170 175

Tyr Cys Asp Phe Glu Val Gln Tyr Glu Ile Leu His Asn Ala Val His
 180 185 190

Ser Trp Leu Gly Gly Thr Gly Gln Tyr Ser Met Ser Thr Leu Glu Tyr
 195 200 205

Ser Ala Phe Asp Pro Val Phe Met Ile His His Ser Ser Leu Asp Arg
 210 215 220

Ile Trp Ile Leu Trp Gln Lys Leu Gln Lys Ile Arg Met Lys Pro Tyr
 225 230 235 240

Tyr Ala Leu Asp Cys Ala Gly Asp Arg Leu Met Lys Asp Pro Leu His
 245 250 255

Pro Phe Asn Tyr Glu Thr Val Asn Glu Asp Glu Phe Thr Arg Ile Asn
 260 265 270

Ser Phe Pro Ser Ile Leu Phe Asp His Tyr Arg Phe Asn Tyr Glu Tyr
 275 280 285

Asp Asn Met Arg Ile Arg Gly Gln Asp Ile His Glu Leu Glu Glu Val
 290 295 300

Ile Gln Glu Leu Arg Asn Lys Asp Arg Ile Phe Ala Gly Phe Val Leu
 305 310 315 320

Ser Gly Leu Arg Ile Ser Ala Thr Val Lys Val Phe Ile His Ser Lys
 325 330 335

Asn Asp Thr Ser His Glu Glu Tyr Ala Gly Glu Phe Ala Val Leu Gly
 340 345 350

Gly Glu Lys Glu Met Pro Trp Ala Tyr Glu Arg Met Leu Lys Leu Asp
 355 360 365

Ile Ser Asp Ala Val His Lys Leu His Val Lys Asp Glu Asp Ile Arg
 370 375 380

Phe Arg Val Val Val Thr Ala Tyr Asn Gly Asp Val Val Thr Thr Arg
 385 390 395 400

Leu Ser Gln Pro Phe Ile Val His Arg Pro Ala His Val Ala His Asp
 405 410 415

Ile Leu Val Ile Pro Val Gly Ala Gly His Asp Leu Pro Pro Lys Val
 420 425 430

Val Val Lys Ser Gly Thr Lys Val Glu Phe Thr Pro Ile Asp Ser Ser
 435 440 445

Val Asn Lys Ala Met Val Glu Leu Gly Ser Tyr Thr Ala Met Ala Lys
 450 455 460

Cys Ile Val Pro Pro Phe Ser Tyr His Gly Phe Glu Leu Asp Lys Val
 465 470 475 480

Tyr Ser Val Asp His Gly Asp Tyr Tyr Ile Ala Ala Gly Thr His Ala
 485 490 495

Leu Cys Glu Gln Asn Leu Arg Leu His Ile His Val Glu His Glu
 500 505 510

<210> 33

<211> 334

<212> PRT

<213> *Haliothis tuberculata*

<400> 33

His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Ile Arg Arg Gly
 1 5 10 15

Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Gln Pro Met Ala His
 20 25 30

Leu Pro Gly Leu Ala Asp Asn Ala Thr Tyr Arg Asp Pro Ile Ser Gly
 35 40 45

Asp Ser Arg His Asn Pro Phe His Asp Val Glu Val Ala Phe Glu Asn
 50 55 60

Gly Arg Thr Glu Arg His Pro Asp Ser Arg Leu Phe Glu Gln Pro Leu
 65 70 75 80

25

Phe Gly Lys His Thr Arg Leu Phe Asp Ser Ile Val Tyr Ala Phe Glu
 85 90 95

Gln Glu Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Met Thr His Asn
 100 105 110

Asn Ile His Ala Trp Ile Gly Gly Glu Lys Tyr Ser Met Ser Ser
 115 120 125

Leu His Tyr Thr Ala Phe Asp Pro Ile Phe Tyr Leu Arg His Ser Asn
 130 135 140

Thr Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg Asn
 145 150 155 160

Arg Pro Tyr Lys Ala His Cys Ala Trp Ser Glu Glu Arg Gln Pro Leu
 165 170 175

Lys Pro Phe Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr Tyr
 180 185 190

Glu Asn Ser Val Pro Thr Asn Val Tyr Asp Tyr Glu Gly Val Leu Gly
 195 200 205

Tyr Thr Tyr Asp Asp Leu Asn Phe Gly Gly Met Asp Leu Gly Gln Leu
 210 215 220

Glu Glu Tyr Ile Gln Arg Gln Arg Asp Arg Thr Phe Ala Gly
 225 230 235 240

Phe Phe Leu Ser His Ile Gly Thr Ser Ala Asn Val Glu Ile Ile Ile
 245 250 255

Asp His Gly Thr Leu His Thr Ser Val Gly Thr Phe Ala Val Leu Gly
 260 265 270

Gly Glu Lys Glu Met Lys Trp Gly Phe Asp Arg Leu Tyr Lys Tyr Glu
 275 280 285

Ile Thr Asp Glu Leu Arg Gln Leu Asn Leu Arg Ala Asp Asp Val Phe
 290 295 300

Ser Ile Ser Val Lys Val Thr Asp Val Asp Gly Ser Glu Leu Ser Ser
 305 310 315 320

Glu Leu Ile Pro Ser Ala Ala Ile Ile Phe Glu Arg Ser His
 325 330

<210> 34

<211> 417

<212> PRT

<213> Haliotis tuberculata

<400> 34

Ile Asp His Gln Asp Pro His His Asp Thr Ile Ile Arg Lys Asn Val
 1 5 10 15

Asp Asn Leu Thr Pro Glu Glu Ile Asn Ser Leu Arg Arg Ala Met Ala
 20 25 30

Asp Leu Gln Ser Asp Lys Thr Ala Gly Gly Phe Gln Gln Ile Ala Ala
 35 40 45

Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Asp Ala Glu Lys Lys
 50 55 60

Phe Ser Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
 65 70 75 80

Leu Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Cys Leu
 85 90 95

Gly Ala Leu Pro Tyr Trp Asp Trp Thr Arg Pro Leu Ser His Leu Pro
 100 105 110

Asp Leu Val Leu Val Ser Ser Arg Thr Thr Pro Met Pro Tyr Ser Thr
 115 120 125

Val Glu Ala Arg Asn Pro Trp Tyr Ser Gly His Ile Asp Thr Val Gly
 130 135 140

Val Asp Thr Thr Arg Ser Val Arg Gln Glu Leu Tyr Glu Ala Pro Gly
 145 150 155 160

Phe Gly His Tyr Thr Gly Val Ala Lys Gln Val Leu Leu Ala Leu Glu
 165 170 175

Gln Asp Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn
 180 185 190

Phe Ile His Ala Leu Val Gly Gly Ser Glu Pro Tyr Gly Met Ala Ser
 195 200 205

Leu Arg Tyr Thr Thr Tyr Asp Pro Ile Phe Tyr Leu His His Ser Asn
 210 215 220

Thr Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly
 225 230 235 240

Lys Pro Tyr Asn Ser Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro
 245 250 255

Leu Gln Pro Phe Gly Leu Thr Asp Glu Ile Asn Pro Asp Asp Glu Thr
 260 265 270

Arg Gln His Ala Val Pro Phe Ser Val Phe Asp Tyr Lys Asn Asn Phe
 275 280 285

Asn Tyr Glu Tyr Asp Thr Leu Asp Phe Asn Gly Leu Ser Ile Ser Gln
 290 295 300

Leu Asp Arg Glu Leu Ser Arg Arg Lys Ser His Asp Arg Val Phe Ala
 305 310 315 320

Gly Phe Leu Leu His Gly Ile Gln Gln Ser Ala Leu Val Lys Phe Phe
 325 330 335
 Val Cys Lys Ser Asp Asp Asp Cys Asp His Tyr Ala Gly Glu Phe Tyr
 340 345 350
 Ile Leu Gly Asp Glu Ala Glu Met Pro Trp Gly Tyr Asp Arg Leu Tyr
 355 360 365
 Lys Tyr Glu Ile Thr Glu Gln Leu Asn Ala Leu Asp Leu His Ile Gly
 370 375 380
 Asp Arg Phe Phe Ile Arg Tyr Glu Ala Phe Asp Leu His Gly Thr Ser
 385 390 395 400
 Leu Gly Ser Asn Ile Phe Pro Lys Pro Ser Val Ile His Asp Glu Gly
 405 410 415
 Ala

<210> 35
 <211> 415
 <212> PRT
 <213> *Haliotis tuberculata*

 <400> 35
 Gly His His Gln Ala Asp Glu Tyr Asp Glu Val Val Thr Ala Ala Ser
 1 5 10 15

 His Ile Arg Lys Asn Leu Lys Asp Leu Ser Lys Gly Glu Val Glu Ser
 20 25 30

 Leu Arg Ser Ala Phe Leu Gln Leu Gln Asn Asp Gly Val Tyr Glu Asn
 35 40 45

 Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Asp Asp Asn Gly Arg
 50 55 60

 Lys Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
 65 70 75 80

 Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95

 Ala Val Ser Val Pro Tyr Trp Asp Trp Thr Glu Thr Phe Thr Glu Leu
 100 105 110

 Pro Ser Leu Ile Ala Glu Ala Thr Tyr Phe Asn Ser Arg Gln Gln Thr
 115 120 125

 Phe Asp Pro Asn Pro Phe Phe Arg Gly Lys Ile Ser Phe Glu Asn Ala
 130 135 140

28

Val Thr Thr Arg Asp Pro Gln Pro Glu Leu Tyr Val Asn Arg Tyr Tyr
 145 150 155 160
 Tyr Gln Asn Val Met Leu Val Phe Glu Gln Asp Asn Tyr Cys Asp Phe
 165 170 175
 Glu Ile Gln Phe Glu Met Val His Asn Val Leu His Ala Trp Leu Gly
 180 185 190
 Gly Arg Ala Thr Tyr Ser Ile Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205
 Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220
 Trp Gln Glu Leu Gln Arg Tyr Arg Lys Lys Pro Tyr Asn Glu Ala Asp
 225 230 235 240
 Cys Ala Ile Asn Leu Met Arg Lys Pro Leu His Pro Phe Asp Asn Ser
 245 250 255
 Asp Leu Asn His Asp Pro Val Thr Phe Lys Tyr Ser Lys Pro Thr Asp
 260 265 270
 Gly Phe Asp Tyr Gln Asn Asn Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285
 Phe Asn His Phe Ser Ile Pro Arg Leu Glu Glu Ile Ile Arg Ile Arg
 290 295 300
 Gln Arg Gln Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320
 Thr Ser Ala Thr Val Glu Ile Phe Val Cys Val Pro Thr Thr Ser Gly
 325 330 335
 Glu Gln Asn Cys Glu Asn Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 340 345 350
 Glu Thr Glu Met Ala Phe His Phe Asp Arg Leu Tyr Arg Phe Asp Ile
 355 360 365
 Ser Glu Thr Leu Arg Asp Leu Gly Ile Gln Leu Asp Ser His Asp Phe
 370 375 380
 Asp Leu Ser Ile Lys Ile Gln Gly Val Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400
 His Ile Leu Pro Glu Pro Ser Leu Ile Phe Val Pro Gly Ser Ser
 405 410 415

<210> 36
 <211> 418
 <212> PRT
 <213> *Haliothis tuberculata*

29

<400> 36

Ser Phe Leu Arg Pro Asp Gly His Ser Asp Asp Ile Leu Val Arg Lys
 1 5 10 15

Glu Val Asn Ser Leu Thr Thr Arg Glu Thr Ala Ser Leu Ile His Ala
 20 25 30

Leu Lys Ser Met Gln Glu Asp His Ser Pro Asp Gly Phe Gln Ala Ile
 35 40 45

Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala Ala
 50 55 60

His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln Trp
 65 70 75 80

His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ala Leu Arg Arg His Gly
 85 90 95

Ala Thr Val Gly Val Pro Tyr Trp Asp Trp Leu Arg Pro Gln Ser His
 100 105 110

Leu Pro Glu Leu Val Thr Met Glu Thr Tyr His Asp Ile Trp Ser Asn
 115 120 125

Arg Asp Phe Pro Asn Pro Phe Tyr Gln Ala Asn Ile Glu Phe Glu Gly
 130 135 140

Glu Asn Ile Thr Thr Glu Arg Glu Val Ile Ala Asp Lys Leu Phe Val
 145 150 155 160

Lys Gly Gly His Val Phe Asp Lys Leu Val Leu Gln Thr Ser His Pro
 165 170 175

Ser Ala Glu Gln Glu Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu Ile
 180 185 190

Leu His Asn Gly Val His Thr Trp Val Gly Gly Ser Arg Thr Tyr Ser
 195 200 205

Ile Gly His Leu His Tyr Ala Phe Tyr Asp Pro Leu Phe Tyr Leu His
 210 215 220

His Phe Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu Gln Glu
 225 230 235 240

Gln Arg Gly Leu Ser Gly Asp Glu Ala His Cys Ala Leu Glu Gln Met
 245 250 255

Arg Glu Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Trp Asn
 260 265 270

Gln Leu Thr Gln Asp Phe Ser Arg Pro Glu Asp Thr Phe Asp Tyr Arg
 275 280 285

Lys Phe Gly Tyr Glu Tyr Asp Asn Leu Glu Phe Leu Gly Met Ser Val
 290 295 300

30

Ala Glu Leu Asp Gln Tyr Ile Ile Glu His Gln Glu Asn Asp Arg Val
305 310 315 320

Phe Ala Gly Phe Leu Leu Ser Gly Phe Gly Gly Ser Ala Ser Val Asn
325 330 335

Phe Gln Val Cys Arg Ala Asp Ser Thr Cys Gln Asp Ala Gly Tyr Phe
340 345 350

Thr Val Leu Gly Gly Ser Ala Glu Met Ala Trp Ala Phe Asp Arg Leu
355 360 365

Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Glu Lys Met His Leu Arg Tyr
370 375 380

Asp Asp Asp Phe Thr Ile Ser Val Ser Leu Thr Ala Asn Asn Gly Thr
385 390 395 400

Val Leu Ser Ser Ser Leu Ile Pro Thr Pro Ser Val Ile Phe Gln Arg
405 410 415

Gly His

<210> 37
<211> 416
<212> PRT
<213> *Haliotis tuberculata*

<400> 37
Arg Asp Ile Asn Thr Arg Ser Met Ser Pro Asn Arg Val Arg Arg Glu
1 5 10 15

Leu Ser Asp Leu Ser Ala Arg Asp Leu Ser Ser Leu Lys Ser Ala Leu
20 25 30

Arg Asp Leu Gln Glu Asp Asp Gly Pro Asn Gly Tyr Gln Ala Leu Ala
35 40 45

Ala Phe His Gly Leu Pro Ala Gly Cys His Asp Ser Arg Gly Asn Glu
50 55 60

Ile Ala Cys Cys Ile His Gly Met Pro Thr Phe Pro Gln Trp His Arg
65 70 75 80

Leu Tyr Thr Leu Gln Leu Glu Met Ala Leu Arg Arg His Gly Ser Ser
85 90 95

Val Ala Ile Pro Tyr Trp Asp Trp Thr Lys Pro Ile Ser Glu Leu Pro
100 105 110

Ser Leu Phe Thr Ser Pro Glu Tyr Tyr Asp Pro Trp His Asp Ala Val
115 120 125

31

Val Asn Asn Pro Phe Ser Lys Gly Phe Val Lys Phe Ala Asn Thr Tyr
130 135 140

Thr Val Arg Asp Pro Gln Glu Met Leu Phe Gln Leu Cys Glu His Gly
145 150 155 160

Glu Ser Ile Leu Tyr Glu Gln Thr Leu Leu Ala Leu Glu Gln Thr Asp
165 170 175

Tyr Cys Asp Phe Glu Val Gln Phe Glu Val Leu His Asn Val Ile His
180 185 190

Tyr Leu Val Gly Gly Arg Gln Thr Tyr Ala Leu Ser Ser Leu His Tyr
195 200 205

Ala Ser Tyr Asp Pro Phe Phe Ile His His Ser Phe Val Asp Lys
210 215 220

Met Trp Val Val Trp Gln Ala Leu Gln Lys Arg Arg Lys Leu Pro Tyr
225 230 235 240

Lys Arg Ala Asp Cys Ala Val Asn Leu Met Thr Lys Pro Met Arg Pro
245 250 255

Phe Asp Ser Asp Met Asn Gln Asn Pro Phe Thr Lys Met His Ala Val
260 265 270

Pro Asn Thr Leu Tyr Asp Tyr Glu Thr Leu Tyr Tyr Ser Tyr Asp Asn
275 280 285

Leu Glu Ile Gly Gly Arg Asn Leu Asp Gln Leu Gln Ala Glu Ile Asp
290 295 300

Arg Ser Arg Ser His Asp Arg Val Phe Ala Gly Phe Leu Leu Arg Gly
305 310 315 320

Ile Gly Thr Ser Ala Asp Val Arg Phe Trp Ile Cys Arg Asn Glu Asn
325 330 335

Asp Cys His Arg Gly Gly Ile Ile Phe Ile Leu Gly Gly Ala Lys Glu
340 345 350

Met Pro Trp Ser Phe Asp Arg Asn Phe Lys Phe Asp Ile Thr His Val
355 360 365

Leu Glu Asn Ala Gly Ile Ser Pro Glu Asp Val Phe Asp Ala Glu Glu
370 375 380

Pro Phe Tyr Ile Lys Val Glu Ile His Ala Val Asn Lys Thr Met Ile
385 390 395 400

Pro Ser Ser Val Ile Pro Ala Pro Thr Ile Ile Tyr Ser Pro Gly Glu
405 410 415

<210> 38
 <211> 402
 <212> PRT
 <213> *Haliotis tuberculata*

<400> 38
 Gly Arg Ala Ala Asp Ser Ala His Ser Ala Asn Ile Ala Gly Ser Gly
 1 5 10 15

Val Arg Lys Asp Val Thr Thr Leu Thr Val Ser Glu Thr Glu Asn Leu
 20 25 30

Arg Gln Ala Leu Gln Gly Val Ile Asp Asp Thr Gly Pro Asn Gly Tyr
 35 40 45

Gln Ala Ile Ala Ser Phe His Gly Ser Pro Pro Met Cys Glu Met Asn
 50 55 60

Gly Arg Lys Val Ala Cys Cys Ala His Gly Met Ala Ser Phe Pro His
 65 70 75 80

Trp His Arg Leu Tyr Val Lys Gln Met Glu Asp Ala Leu Ala Asp His
 85 90 95

Gly Ser His Ile Gly Ile Pro Tyr Trp Asp Trp Thr Thr Ala Phe Thr
 100 105 110

Glu Leu Pro Ala Leu Val Thr Asp Ser Glu Asn Asn Pro Phe His Glu
 115 120 125

Gly Arg Ile Asp His Leu Gly Val Thr Thr Ser Arg Ser Pro Arg Asp
 130 135 140

Met Leu Phe Asn Asp Pro Glu Gln Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160

Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Tyr Cys Gln Phe Glu Val
 165 170 175

Gln Phe Glu Leu Thr His Asn Ala Ile His Ser Trp Thr Gly Gly Arg
 180 185 190

Ser Pro Tyr Gly Met Ser Thr Leu Glu Phe Thr Ala Tyr Asp Pro Leu
 195 200 205

Phe Trp Leu His His Ser Asn Thr Asp Arg Ile Trp Ala Val Trp Gln
 210 215 220

Ala Leu Gln Lys Tyr Arg Gly Leu Pro Tyr Asn Glu Ala His Cys Glu
 225 230 235 240

Ile Gln Val Leu Lys Gln Pro Leu Arg Pro Phe Asn Asp Asp Ile Asn
 245 250 255

His Asn Pro Ile Thr Lys Thr Asn Ala Arg Pro Ile Asp Ser Phe Asp
 260 265 270

Tyr Glu Arg Phe Asn Tyr Gln Tyr Asp Thr Leu Ser Phe His Gly Lys
 275 280 285
 Ser Ile Pro Glu Leu Asn Asp Leu Leu Glu Glu Arg Lys Arg Glu Glu
 290 295 300
 Arg Thr Phe Ala Ala Phe Leu Leu Arg Gly Ile Gly Cys Ser Ala Asp
 305 310 315 320
 Val Val Phe Asp Ile Cys Arg Pro Asn Gly Asp Cys Val Phe Ala Gly
 325 330 335
 Thr Phe Ala Val Leu Gly Gly Glu Leu Glu Met Pro Trp Ser Phe Asp
 340 345 350
 Arg Leu Phe Arg Tyr Asp Ile Thr Arg Val Met Asn Gln Leu His Leu
 355 360 365
 Gln Tyr Asp Ser Asp Phe Ser Phe Arg Val Lys Leu Val Ala Thr Asn
 370 375 380
 Gly Thr Glu Leu Ser Ser Asp Leu Leu Lys Ser Pro Thr Ile Glu His
 385 390 395 400
 Glu Leu

<210> 39
 <211> 515
 <212> PRT
 <213> *Haliotis tuberculata*

<400> 39
 Gly Ala His Arg Gly Pro Val Glu Glu Thr Glu Val Thr Arg Gln His
 1 5 10 15
 Thr Asp Gly Asn Ala His Phe His Arg Lys Glu Val Asp Ser Leu Ser
 20 25 30
 Leu Asp Glu Ala Asn Asn Leu Lys Asn Ala Leu Tyr Lys Leu Gln Asn
 35 40 45
 Asp His Ser Leu Thr Gly Tyr Glu Ala Ile Ser Gly Tyr His Gly Tyr
 50 55 60
 Pro Asn Leu Cys Pro Glu Glu Gly Asp Asp Lys Ile Pro Leu Leu Arg
 65 70 75 80
 Pro Arg Met Gly Ile Phe Pro Tyr Trp His Arg Leu Leu Thr Ile Gln
 85 90 95
 Leu Glu Arg Ala Leu Glu His Asn Gly Ala Leu Leu Gly Val Pro Tyr
 100 105 110

34

Trp Asp Trp Asn Lys Asp Leu Ser Ser Leu Pro Ala Phe Phe Ser Asp
 115 120 125

Ser Ser Asn Asn Asn Pro Tyr Phe Lys Tyr His Ile Ala Gly Val Gly
 130 135 140

His Asp Thr Val Arg Glu Pro Thr Ser Leu Ile Tyr Asn Gln Pro Gln
 145 150 155 160

Ile His Gly Tyr Asp Tyr Leu Tyr Tyr Leu Ala Leu Thr Thr Leu Glu
 165 170 175

Glu Asn Asn Tyr Trp Asp Phe Glu Val Gln Tyr Glu Ile Leu His Asn
 180 185 190

Ala Val His Ser Trp Leu Gly Gly Ser Gln Lys Tyr Ser Met Ser Thr
 195 200 205

Leu Glu Tyr Ser Ala Phe Asp Pro Val Phe Met Ile Leu His Ser Gly
 210 215 220

Leu Asp Arg Leu Trp Ile Ile Trp Gln Glu Leu Gln Lys Ile Arg Arg
 225 230 235 240

Lys Pro Tyr Asn Phe Ala Lys Cys Ala Tyr His Met Met Glu Glu Pro
 245 250 255

Leu Ala Pro Phe Ser Tyr Pro Ser Ile Asn Gln Asp Glu Phe Thr Arg
 260 265 270

Ala Asn Ser Lys Pro Ser Thr Val Phe Asp Ser His Lys Phe Gly Tyr
 275 280 285

His Tyr Asp Asn Leu Asn Val Arg Gly His Ser Ile Gln Glu Leu Asn
 290 295 300

Thr Ile Ile Asn Asp Leu Arg Asn Thr Asp Arg Ile Tyr Ala Gly Phe
 305 310 315 320

Val Leu Ser Gly Ile Gly Thr Ser Ala Ser Val Lys Ile Tyr Leu Arg
 325 330 335

Thr Asp Asp Asn Asp Glu Glu Val Gly Thr Phe Thr Val Leu Gly Gly
 340 345 350

Glu Arg Glu Met Pro Trp Ala Tyr Glu Arg Val Phe Lys Tyr Asp Ile
 355 360 365

Thr Glu Val Ala Asp Arg Leu Lys Ile Lys Leu Trp Gly His Pro Leu
 370 375 380

Thr Ser Gly Thr Gly Asp His Ile Leu Thr Asn Gly Ile Gly Gly Lys
 385 390 395 400

Gln Glu Pro Thr Gln Ile Leu Ser Ser Ser Thr Asp Leu Pro Ile Met
 405 410 415

35

Thr Thr Met Phe Leu Leu Ser Gln Xaa Gly Arg Asn Leu His Ile Pro
 420 425 430

Pro Lys Val Val Val Lys Lys Gly Thr Arg Ile Glu Phe His Pro Val
 435 440 445

Asp Asp Ser Val Thr Arg Pro Val Val Asp Leu Gly Ser Tyr Thr Ala
 450 455 460

Leu Phe Asn Cys Val Val Pro Pro Phe Thr Tyr His Gly Phe Glu Leu
 465 470 475 480

Asn His Val Tyr Ser Val Lys Pro Gly Asp Tyr Tyr Val Thr Gly Pro
 485 490 495

Thr Arg Asp Leu Cys Gln Asn Ala Asp Val Arg Ile His Ile His Val
 500 505 510

Glu Asp Glu
 515

<210> 40

<211> 322

<212> PRT

<213> Megathura crenulata

<400> 40

Gly Leu Pro Tyr Trp Asp Trp Thr Glu Pro Met Thr His Ile Pro Gly
 1 5 10 15

Leu Ala Gly Asn Lys Thr Tyr Val Asp Ser His Gly Ala Ser His Thr
 20 25 30

Asn Pro Phe His Ser Ser Val Ile Ala Phe Glu Glu Asn Ala Pro His
 35 40 45

Thr Lys Arg Gln Ile Asp Gln Arg Leu Phe Lys Pro Ala Thr Phe Gly
 50 55 60

His His Thr Asp Leu Phe Asn Gln Ile Leu Tyr Ala Phe Glu Gln Glu
 65 70 75 80

Asp Tyr Cys Asp Phe Glu Val Gln Phe Glu Ile Thr His Asn Thr Ile
 85 90 95

His Ala Trp Thr Gly Gly Ser Glu His Phe Ser Met Ser Ser Leu His
 100 105 110

Tyr Thr Ala Phe Asp Pro Leu Phe Tyr Phe His His Ser Asn Val Asp
 115 120 125

Arg Leu Trp Ala Val Trp Gln Ala Leu Gln Met Arg Arg His Lys Pro
 130 135 140

Tyr Arg Ala His Cys Ala Ile Ser Leu Glu His Met His Leu Lys Pro
 145 150 155 160

Phe Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr His Ala Asn
 165 170 175
 Ala Met Pro Asn Lys Ile Tyr Asp Tyr Glu Asn Val Leu His Tyr Thr
 180 185 190
 Tyr Glu Asp Leu Thr Phe Gly Gly Ile Ser Leu Glu Asn Ile Glu Lys
 195 200 205
 Met Ile His Glu Asn Gln Gln Glu Asp Arg Ile Tyr Ala Gly Phe Leu
 210 215 220
 Leu Ala Gly Ile Arg Thr Ser Ala Asn Val Asp Ile Phe Ile Lys Thr
 225 230 235 240
 Thr Asp Ser Val Gln His Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 245 250 255
 Ser Lys Glu Met Lys Trp Gly Phe Asp Arg Val Phe Lys Phe Asp Ile
 260 265 270
 Thr His Val Leu Lys Asp Leu Asp Leu Thr Ala Asp Gly Asp Phe Glu
 275 280 285
 Val Thr Val Asp Ile Thr Glu Val Asp Gly Thr Lys Leu Ala Ser Ser
 290 295 300
 Leu Ile Pro His Ala Ser Val Ile Arg Glu His Ala Arg Gly Lys Leu
 305 310 315 320
 Asn Arg

<210> 41
 <211> 414
 <212> PRT
 <213> Megathura crenulata

<400> 41
 Val Lys Phe Asp Lys Val Pro Arg Ser Arg Leu Ile Arg Lys Asn Val
 1 5 10 15
 Asp Arg Leu Ser Pro Glu Glu Met Asn Glu Leu Arg Lys Ala Leu Ala
 20 25 30
 Leu Leu Lys Glu Asp Lys Ser Ala Gly Gly Phe Gln Gln Leu Gly Ala
 35 40 45
 Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Ser Lys Lys
 50 55 60
 Phe Ala Cys Cys Val His Gly Met Ser Val Phe Pro His Trp His Arg
 65 70 75 80

37

Leu Leu Thr Val Gln Ser Glu Asn Ala Leu Arg Arg His Gly Tyr Asp
 85 90 95

Gly Ala Leu Pro Tyr Trp Asp Trp Thr Ser Pro Leu Asn His Leu Pro
 100 105 110

Glu Leu Ala Asp His Glu Lys Tyr Val Asp Pro Glu Asp Gly Val Glu
 115 120 125

Lys His Asn Pro Trp Phe Asp Gly His Ile Asp Thr Val Asp Lys Thr
 130 135 140

Thr Thr Arg Ser Val Gln Asn Lys Leu Phe Glu Gln Pro Glu Phe Gly
 145 150 155 160

His Tyr Thr Ser Ile Ala Lys Gln Val Leu Leu Ala Leu Glu Gln Asp
 165 170 175

Asn Phe Cys Asp Phe Glu Ile Gln Tyr Glu Ile Ala His Asn Tyr Ile
 180 185 190

His Ala Leu Val Gly Gly Ala Gln Pro Tyr Gly Met Ala Ser Leu Arg
 195 200 205

Tyr Thr Ala Phe Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr Asp
 210 215 220

Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro
 225 230 235 240

Tyr Asn Val Ala Asn Cys Ala Val Thr Ser Met Arg Glu Pro Leu Gln
 245 250 255

Pro Phe Gly Leu Ser Ala Asn Ile Asn Thr Asp His Val Thr Lys Glu
 260 265 270

His Ser Val Pro Phe Asn Val Phe Asp Tyr Lys Thr Asn Phe Asn Tyr
 275 280 285

Glu Tyr Asp Thr Leu Glu Phe Asn Gly Leu Ser Ile Ser Gln Leu Asn
 290 295 300

Lys Lys Leu Glu Ala Ile Lys Ser Gln Asp Arg Phe Phe Ala Gly Phe
 305 310 315 320

Leu Leu Ser Gly Phe Lys Lys Ser Ser Leu Val Lys Phe Asn Ile Cys
 325 330 335

Thr Asp Ser Ser Asn Cys His Pro Ala Gly Glu Phe Tyr Leu Leu Gly
 340 345 350

Asp Glu Asn Glu Met Pro Trp Ala Tyr Asp Arg Val Phe Lys Tyr Asp
 355 360 365

Ile Thr Glu Lys Leu His Asp Leu Lys Leu His Ala Glu Asp His Phe
 370 375 380

38

Tyr Ile Asp Tyr Glu Val Phe Asp Leu Lys Pro Ala Ser Leu Gly Lys
 385 390 395 400

Asp Leu Phe Lys Gln Pro Ser Val Ile His Glu Pro Arg Ile
 405 410

<210> 42
 <211> 411
 <212> PRT
 <213> Megathura crenulata

<400> 42
 Gly His His Glu Gly Glu Val Tyr Gln Ala Glu Val Thr Ser Ala Asn
 1 5 10 15

Arg Ile Arg Lys Asn Ile Glu Asn Leu Ser Leu Gly Glu Leu Glu Ser
 20 25 30

Leu Arg Ala Ala Phe Leu Glu Ile Glu Asn Asp Gly Thr Tyr Glu Ser
 35 40 45

Ile Ala Lys Phe His Gly Ser Pro Gly Leu Cys Gln Leu Asn Gly Asn
 50 55 60

Pro Ile Ser Cys Cys Val His Gly Met Pro Thr Phe Pro His Trp His
 65 70 75 80

Arg Leu Tyr Val Val Val Glu Asn Ala Leu Leu Lys Lys Gly Ser
 85 90 95

Ser Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Arg Ile Glu His Leu
 100 105 110

Pro His Leu Ile Ser Asp Ala Thr Tyr Tyr Asn Ser Arg Gln His His
 115 120 125

Tyr Glu Thr Asn Pro Phe His His Gly Lys Ile Thr His Glu Asn Glu
 130 135 140

Ile Thr Thr Arg Asp Pro Lys Asp Ser Leu Phe His Ser Asp Tyr Phe
 145 150 155 160

Tyr Glu Gln Val Leu Tyr Ala Leu Glu Gln Asp Asn Phe Cys Asp Phe
 165 170 175

Glu Ile Gln Leu Glu Ile Leu His Asn Ala Leu His Ser Leu Leu Gly
 180 185 190

Gly Lys Gly Lys Tyr Ser Met Ser Asn Leu Asp Tyr Ala Ala Phe Asp
 195 200 205

Pro Val Phe Phe Leu His His Ala Thr Thr Asp Arg Ile Trp Ala Ile
 210 215 220 240

Trp Gln Asp Leu Gln Arg Phe Arg Lys Arg Pro Tyr Arg Glu Ala Asn
 225 230 235 240

Cys Ala Ile Gln Leu Met His Thr Pro Leu Gln Pro Phe Asp Lys Ser
 245 250 255
 Asp Asn Asn Asp Glu Ala Thr Lys Thr His Ala Thr Pro His Asp Gly
 260 265 270
 Phe Glu Tyr Gln Asn Ser Phe Gly Tyr Ala Tyr Asp Asn Leu Glu Leu
 275 280 285
 Asn His Tyr Ser Ile Pro Gln Leu Asp His Met Leu Gln Glu Arg Lys
 290 295 300
 Arg His Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly Thr
 305 310 315 320
 Ser Ala Asp Gly His Val Phe Val Cys Leu Pro Thr Gly Glu His Thr
 325 330 335
 Lys Asp Cys Ser His Glu Ala Gly Met Phe Ser Ile Leu Gly Gly Gln
 340 345 350
 Thr Glu Met Ser Phe Val Phe Asp Arg Leu Tyr Lys Leu Asp Ile Thr
 355 360 365
 Lys Ala Leu Lys Lys Asn Gly Val His Leu Gln Gly Asp Phe Asp Leu
 370 375 380
 Glu Ile Glu Ile Thr Ala Val Asn Gly Ser His Leu Asp Ser His Val
 385 390 395 400
 Ile His Ser Pro Thr Ile Leu Phe Glu Ala Gly
 405 410

<210> 43
 <211> 111
 <212> PRT
 <213> Megathura crenulata

<400> 43
 Asp Ser Ala His Thr Asp Asp Gly His Thr Glu Pro Val Met Ile Arg
 1 5 10 15
 Lys Asp Ile Thr Gln Leu Asp Lys Arg Gln Gln Leu Ser Leu Val Lys
 20 25 30
 Ala Leu Glu Ser Met Lys Ala Asp His Ser Ser Asp Gly Phe Gln Ala
 35 40 45
 Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ala Ala
 50 55 60
 Ser Lys Arg Phe Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln
 65 70 75 80

40

Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ser Leu Arg Lys His
 85 90 95

Gly Ala Val Val Gly Leu Pro Tyr Trp Asp Trp Thr Leu Pro Arg
 100 105 110

<210> 44
 <211> 317
 <212> PRT
 <213> Megathura crenulata

<400> 44

Gly Leu Pro Tyr Trp Asp Trp Thr Met Pro Met Ser His Leu Pro Glu
 1 5 10 15

Leu Ala Thr Ser Glu Thr Tyr Leu Asp Pro Val Thr Gly Glu Thr Lys
 20 25 30

Asn Asn Pro Phe His His Ala Gln Val Ala Phe Glu Asn Gly Val Thr
 35 40 45

Ser Arg Asn Pro Asp Ala Lys Leu Phe Met Lys Pro Thr Tyr Gly Asp
 50 55 60

His Thr Tyr Leu Phe Asp Ser Met Ile Tyr Ala Phe Glu Gln Glu Asp
 65 70 75 80

Phe Cys Asp Phe Glu Val Gln Tyr Glu Leu Thr His Asn Ala Ile His
 85 90 95

Ala Trp Val Gly Gly Ser Glu Lys Tyr Ser Met Ser Ser Leu His Tyr
 100 105 110

Thr Ala Phe Asp Pro Ile Phe Tyr Leu His His Ser Asn Val Asp Arg
 115 120 125

Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg Gly Lys Ser Tyr
 130 135 140

Lys Ala His Cys Ala Ser Ser Gln Glu Arg Glu Pro Leu Lys Pro Phe
 145 150 155 160

Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr Tyr His Asn Ser
 165 170 175

Val Pro Thr Asn Val Tyr Asp Tyr Val Gly Val Leu His Tyr Arg Tyr
 180 185 190

Asp Asp Leu Gln Phe Gly Gly Met Thr Met Ser Glu Leu Glu Glu Tyr
 195 200 205

Ile His Lys Gln Thr Gln His Asp Arg Thr Phe Ala Gly Phe Phe Leu
 210 215 220

Ser Tyr Ile Gly Thr Ser Ala Ser Val Asp Ile Phe Ile Asn Arg Glu
 225 230 235 240

41

Gly His Asp Lys Tyr Lys Val Gly Ser Phe Val Val Leu Gly Gly Ser
 245 250 255

Lys Glu Met Lys Trp Gly Phe Asp Arg Met Tyr Lys Tyr Glu Ile Thr
 260 265 270

Glu Ala Leu Lys Thr Leu Asn Val Ala Val Asp Asp Gly Phe Ser Ile
 275 280 285

Thr Val Glu Ile Thr Asp Val Asp Gly Ser Pro Pro Ser Ala Asp Leu
 290 295 300

Ile Pro Pro Pro Ala Ile Ile Phe Glu Arg Gly His Ala
 305 310 315

<210> 45

<211> 411

<212> PRT

<213> Megathura crenulata

<400> 45

Asp Ala Lys Asp Phe Gly His Ser Arg Lys Ile Arg Lys Ala Val Asp
 1 5 10 15

Ser Leu Thr Val Glu Glu Gln Thr Ser Leu Arg Arg Ala Met Ala Asp
 20 25 30

Leu Gln Asp Asp Lys Thr Ser Gly Gly Phe Gln Gln Ile Ala Ala Phe
 35 40 45

His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Glu Lys Lys Phe
 50 55 60

Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg Leu
 65 70 75 80

Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Phe Thr Gly
 85 90 95

Gly Leu Pro Tyr Trp Asp Trp Thr Arg Ser Met Ser Ala Leu Pro His
 100 105 110

Phe Val Ala Asp Pro Thr Tyr Asn Asp Ala Ile Ser Ser Gln Glu Glu
 115 120 125

Asp Asn Pro Trp His His Gly His Ile Asp Ser Val Gly His Asp Thr
 130 135 140

Thr Arg Asp Val Arg Asp Asp Leu Tyr Gln Ser Pro Gly Phe Gly His
 145 150 155 160

Tyr Thr Asp Ile Ala Gln Gln Val Leu Leu Ala Phe Glu Gln Asp Ser
 165 170 175

42

Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn Phe Ile His
 180 185 190

Ala Leu Ile Gly Gly Asn Glu Pro Tyr Ser Met Ser Ser Leu Arg Tyr
 195 200 205

Thr Thr Tyr Asp Pro Ile Phe Phe Leu His His Ser Ser Thr Asp Arg
 210 215 220

Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro Tyr
 225 230 235 240

Asn Thr Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro Leu Gln Pro
 245 250 255

Phe Gly Leu Asp Ser Val Ile Asn Pro Asp Asp Glu Thr Arg Glu His
 260 265 270

Ser Val Pro Phe Arg Val Phe Asp Tyr Lys Asn Asn Phe Asp Tyr Glu
 275 280 285

Tyr Glu Ser Leu Ala Phe Asn Gly Leu Ser Ile Ala Gln Leu Asp Arg
 290 295 300

Glu Leu Gln Arg Arg Lys Ser His Asp Arg Val Phe Ala Gly Phe Leu
 305 310 315 320

Leu His Glu Ile Gly Gln Ser Ala Lys His Asn Val Ser Asp Cys Asp
 325 330 335

His Tyr Ala Gly Glu Phe Tyr Ile Leu Gly Asp Glu Ala Glu Met Pro
 340 345 350

Trp Arg Tyr Asp Arg Val Tyr Lys Tyr Glu Ile Thr Gln Gln Leu His
 355 360 365

Asp Leu Asp Leu His Val Gly Asp Asn Phe Phe Leu Lys Tyr Glu Ala
 370 375 380

Phe Asp Leu Asn Gly Gly Ser Leu Gly Gly Ser Ile Phe Ser Gln Pro
 385 390 395 400

Ser Val Ile Phe Glu Pro Ala Ala Gly Met Phe
 405 410

<210> 46

<211> 109

<212> PRT

<213> Megathura crenulata

<400> 46

Gly Ser His Gln Ala Asp Glu Tyr Arg Glu Ala Val Thr Ser Ala Ser
 1 5 10 15

His Ile Arg Lys Asn Ile Arg Asp Leu Ser Glu Gly Glu Ile Glu Ser
 20 25 30

Ile Arg Ser Ala Phe Leu Gln Ile Gln Lys Glu Gly Ile Tyr Glu Asn
 35 40 45
 Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Glu His Asp Gly His
 50 55 60
 Pro Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro His Trp His
 65 70 75 80
 Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95
 Ala Val Ala Val Pro Tyr Trp Asp Trp Thr Leu Pro Arg
 100 105

<210> 47
 <211> 329
 <212> PRT
 <213> Megathura crenulata

<400> 47
 Met Ala Val Phe Pro His Trp His Arg Leu Phe Val Lys Gln Met Glu
 1 5 10 15
 Asp Ala Leu Ala Ala His Gly Ala His Ile Gly Ile Pro Tyr Trp Asp
 20 25 30
 Trp Thr Ser Ala Phe Ser His Leu Pro Ala Leu Val Thr Asp His Glu
 35 40 45
 Asn Asn Pro Phe His His Gly His Ile Gly His Leu Asn Val Asp Thr
 50 55 60
 Ser Arg Ser Pro Arg Asp Met Leu Phe Asn Asp Pro Glu Gln Gly Ser
 65 70 75 80
 Glu Ser Phe Phe Tyr Arg Gln Val Leu Leu Thr Leu Glu Gln Thr Asp
 85 90 95
 Phe Cys Gln Phe Glu Val Gln Phe Glu Leu Thr His Asn Ala Ile His
 100 105 110
 Ser Trp Thr Gly Gly His Thr Pro Tyr Gly Met Ser Ser Leu Glu Tyr
 115 120 125
 Thr Ala Tyr Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr Asp Arg
 130 135 140
 Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Leu Pro Tyr
 145 150 155 160
 Asn Ala Ala His Cys Asp Ile Gln Val Leu Lys Gln Pro Leu Lys Pro
 165 170 175

44

Phe Ser Glu Ser Arg Asn Pro Asn Pro Val Thr Arg Ala Asn Ser Arg
 180 185 190
 Ala Val Asp Ser Phe Asp Tyr Glu Lys Phe Asn Tyr Gln Tyr Asp Thr
 195 200 205
 Leu Thr Phe His Gly Leu Ser Ile Pro Glu Leu Asp Ala Met Leu Gln
 210 215 220
 Glu Arg Lys Lys Glu Glu Arg Thr Phe Ala Ala Phe Leu Leu His Gly
 225 230 235 240
 Phe Gly Ala Ser Ala Asp Val Ser Phe Asp Val Cys Thr Pro Asp Gly
 245 250 255
 His Cys Ala Phe Ala Gly Thr Phe Ala Val Leu Gly Gly Glu Leu Glu
 260 265 270
 Met Pro Trp Ser Phe Glu Arg Leu Phe Arg Tyr Asp Ile Thr Lys Val
 275 280 285
 Leu Lys Gln Met Asn Leu His Tyr Asp Ser Glu Phe His Phe Glu Leu
 290 295 300
 Lys Ile Val Gly Thr Asp Gly Thr Glu Leu Pro Ser Asp Arg Ile Lys
 305 310 315 320
 Ser Pro Thr Ile Glu His His Gly Gly
 325

<210> 48
 <211> 103
 <212> PRT
 <213> Megathura crenulata

<400> 48
 Gly His Asp His Ser Glu Arg His Asp Gly Phe Phe Arg Lys Glu Val
 1 5 10 15
 Gly Ser Leu Ser Leu Asp Glu Ala Asn Asp Leu Lys Asn Ala Leu Tyr
 20 25 30
 Lys Leu Gln Asn Asp Gln Gly Pro Asn Gly Tyr Glu Ser Ile Ala Gly
 35 40 45
 Tyr His Gly Tyr Pro Phe Leu Cys Pro Glu His Gly Glu Asp Gln Tyr
 50 55 60
 Ala Cys Cys Val His Gly Met Pro Val Phe Pro His Trp His Arg Leu
 65 70 75 80
 His Thr Ile Gln Phe Glu Arg Ala Leu Lys Glu His Gly Ser His Leu
 85 90 95
 Gly Leu Pro Tyr Trp Asp Trp
 100

<210> 49
 <211> 1269
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 49
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 acgtcactgc atctacaggg cctctgagtt tcgaagacat aacatcttac catgcccac 180
 cagcgtcgtg tgactacaag ggacggaaga tcgcctgctg tggccacggatgcccagg 240
 tcccccttctg gcacaggggca tatgtcgcc aagccgagcg ggcactgttg tccaaacgga 300
 agactgtcggtt aatgccttac tgggactgga cgcaaacgct gactcaacttacatctctt 360
 tgactgaacc catctacatt gacagtaaaag gtggaaaggc tcaaaaccaac tactggtacc 420
 gcggcggagat agcgttcatc aataagaaga ctgcgcgagc tggatgatgat cgcctattcg 480
 agaagggtgga gcctggtcac tacacacatc ttatggagac tgccttcgac gctctcgac 540
 aggacgaatt ctgtaaattt gaaatccagt tcgagttggc tcataatgtatccattact 600
 tgggtggcg taaatttggaa tattcaatgt caaacttggaa atacaccccttacgacccca 660
 tcttcttcccttccaccactcc aacgttgcaccgccttccatctggcag cgtcttcagg 720
 aactgcgagg aaagaatcccc aatgcaatgg actgtgcaca tgaactcgctt caccagcaac 780
 tccaaacccttccaccacttccaccactcc aacaggggac agcaatccag tccagcttacaaaggaccac tcgacacccctt 840
 ctgaccccttcttccaccactcc aacttggat acagctacga cagctttaaac ctgaatggaa 900
 tgacgcccaga acagctgaaa acagaacttag acgaacgcca ctccaaagaa cgtgcgtttt 960
 caagcttccg actcagttgc ttgggggtt ctgccaacgt tggatgtctat gcatgtgtcc 1020
 ctgatgtatc tccacgcagt gatgactact gcgagaaagc aggccacttc ttcatctt 1080
 ggggtcaaaag cgaaatcccg tggagattcttccaccacttcttcatgtat gtaactgaag 1140
 cggtacatca ccttggagtc ccgcttaagt gccactacta tggaaaaca gaactcttca 1200
 gctgtgaatgg cacagcaactt tcacctgtatc ttcttcctca accaactgtt gcttaccgac 1260
 ctggggaaag 1269

<210> 50
 <211> 569
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 50
 ggcttccgt actgggactg gacgcagcat ctgactcaac tcccaagatct ggtgtcagac 60
 ccccttggggatcggaggaggaaaag gcccatttgcac acgcatttgcac tcgtggaaac 120
 atcaagtttgcac agaataagaa gactgcaaga gctgttgcac atcccttttgcac 180
 ggaccaggaggaggaaaatcccg actcttgcac ggaattctgc atgccttgcac acaggatgaa 240
 ttctgcacttgcac gtttgcac gcttgcac gtcacaaacgcttccacta cctgggttgc 300
 gggccgtcaca cgtactccatgttcatcttgcac gatgttgcac ctccttgcac ccccttctt 360
 tcctccatca ctccaaacaccggaccgcacccatcttgcac gggacgttgc acaggatgac 420
 agaggaaaagg accccaaacac cggccactgc gcacacaacc tcatccatgc gcccattggaa 480
 ccgttccgtc gggactcgaa cccttgcac ctcaccaggaaaactccaa accaatttgcac 540
 agctttgatttgcac gtttgcac 569

<210> 51
 <211> 1246
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 51
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 catctgacac gcgaggaggat gtacgagctg cgcagagcttgc tggagatggatccaggccgac 120
 acatccgttgcac ggttgcac gtttgcac gtttgcac 180

46

| | | | | | | |
|------------|-------------|-------------|-------------|-------------|-------------|------|
| tcccccgagg | ccacaatag | gttcgcctgt | tgcatccacg | gcatggcgcac | attccctcat | 240 |
| tggcacagac | tgttcgtcac | ccaggtggaa | gatgctctga | tcagggcgagg | atcgccata | 300 |
| ggggtccct | actgggactg | gactcagcc | atggcgcate | tcccaggact | tgcagacaac | 360 |
| gccacctata | gagatcccat | cagcggggac | agcagacaca | acccttcca | cgatgttga | 420 |
| gttgccttg | aaaatggacg | tacagaacgt | cacccagata | gtagattgtt | tgaacaacct | 480 |
| ttatttgca | aacatacgcg | tctcttcgac | agtatagtct | atgttttga | gcagggaggac | 540 |
| ttctgcgatt | ttgaagttca | atttgagatg | accctataata | atattcacgc | ctggatttgtt | 600 |
| ggcggcgaga | agtattccat | gtcttctcta | cactacacag | ccttcgaccc | tatcttctac | 660 |
| cttcgtcact | ccaaacactga | ccggctctgg | gcaatttggc | aagcgttgc | gatacgaaga | 720 |
| aacaggcctt | acaaggctca | tttgcttgg | tctgaggaac | gccagcctct | caaaccttct | 780 |
| gccttcagtt | ccccactgaa | caacaacgaa | aaaacactcg | aaaactcggt | gcccaccaac | 840 |
| gtttacgact | acgaaggagt | ctttggctat | acttatgtatg | acctcaattt | cgggggcatg | 900 |
| gaccctggtc | agcttgagga | atacatccag | aggcagagac | agagagacag | gacccttgc | 960 |
| ggtttcttc | tgtcacatat | ttgttacatca | gcaatgttg | aaatcattat | agaccatggg | 1020 |
| actcttcata | cctccgtggg | cacgtttgc | gttcttggcg | gagagaagga | gatgaaatgg | 1080 |
| ggatttgacc | gtttgtacaa | atatgagatt | acagatgaac | tgaggcaact | taatctccgt | 1140 |
| gctgtatgt | ttttcagcat | ctctgttaaa | gtaactgtatg | ttgtatggcag | tgagctgtcc | 1200 |
| tctgaactca | tcccatactgc | tgctatcattc | ttcgaacgaa | gccata | | 1246 |

<210> 52

<211> 1242

<212> DNA

<213> *Haliotis tuberculata*

<400> 52

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|-------------|-------------|-------------|--------------|-------------|------------|---------|------------|-------|------|
| gtcaccatca | ggctgacgag | tacgacgaa | ttgttaactgc | tgcaagccac | atcaga | aaaga | 60 | | |
| atttaaaaga | tctgtcaaag | ggagaagtag | agagccctaa | gtctgccttc | ctgc | aaacttc | 120 | | |
| agaacgcacgg | agtcttatgag | aatattgcca | agttccacgg | caagcctggg | ttgtgtat | gatg | 180 | | |
| ataaacggtcg | caaggttgcc | tgttgtgtcc | atggaaatgccc | cacccccc | cagtgg | caca | 240 | | |
| ggctctatgt | cctccagggt | gagaatgcct | tgctggagag | aggatctgccc | gtctctgtgc | | 300 | | |
| catactggaa | ctggactgaa | acatttacag | agctgcccattc | tttgattgt | gagg | gctac | 360 | | |
| atttcaat | ccgtcaacaa | acgtttgacc | ctaattccccc | cttcagagg | aaaat | caagt | 420 | | |
| ttgagaatgc | tgttacaaca | cgtgatcccc | agcctgagct | gtacgtaac | agg | tactact | 480 | | |
| accaaaaacgt | catgttggtt | tttgaacagg | acaactactg | cgacttcgag | atacag | tttg | 540 | | |
| agatggttca | caatgttctc | catgcttggc | tttgggttggaa | agactacttat | tctatttctt | | 600 | | |
| ctcttgattt | ttctgcattc | gaccctgtgt | ttttcccttc | ccatgcgaac | acagat | agat | 660 | | |
| tgtggggccat | ctggcaggag | ctgcagagg | acagggaa | gcatacaat | gaag | ccggat | 720 | | |
| gtgccccat | ccta | atgcgc | aaacccctac | atcccttgc | caacagt | gtat | 780 | | |
| atccctgttac | ctttaaaatc | tcaaaacccca | ctgtatggctt | tgactaccag | aaca | actttg | 840 | | |
| gatacaagta | tgacaaac | ctt | gatgttcaat | atttcgtat | tcccagg | gtt | 900 | | |
| ttcgttattt | acaacgt | caa | gatgtgtgt | ttgcaggatt | cctcc | ttcac | aacattgg | 960 | |
| catccgcac | tgtt | gagata | tgcgtctgt | tcccttaccac | cagcgg | ttag | caaaaactgt | 1020 | |
| aaaacaaagc | cgga | acattt | gcccgtactcg | gaggagaa | agagat | ggcg | tttcatttt | 1080 | |
| acagactcta | cagg | tttgac | atcagt | gaaa | cact | gagg | cata | 1140 | |
| gccatgactt | tgac | ctc | agat | tttt | aagg | ataaa | tggtat | cttgc | 1200 |
| acatcc | tc | ccat | cc | ttgat | tttt | ttc | aa | | 1242 |

<210> 53

<211> 1257

<212> DNA

<213> *Haliotis tuberculata*

<400> 53

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gcctgacaaac caggagact gcatactctga tccatgtctt gaaaagtatg caggaagacc 120
attcacctga cgggttccaa gccattgcct ctttccatgc tctgccccca ctctgcctt 180
caccatctgc agctcaccgt tatgttgcgt gtgtccacgg catggctaca tttccccagt 240
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47

ggcacagatt gtacactgta cagttccagg atgcactgag gagacatgga gctacggtag 300
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 agacataccca tggatattgg agtaacagag atttcccaa tccttctac caagccaata 420
 ttgagttga aggagaaaac attacaacag agagagaagt cattgcagac aaacttttg 480
 tcaaagggtgg acacgtttt gataaactgg ttcttcaaac aaggcatctt agcgtgagc 540
 aggaaaaacta ctgtgacttt gagattcagt ttgaaatct tcacaacggc gttcacacgt 600
 gggtcggagg cagtcgtacc tactctatcg gacatcttca ttacgcattc tacgaccctc 660
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 tgcacccctcg atatgatgat gacttcacaa tctctgtcag tctgaccggc aacaacggaa 1200
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<210> 54

<211> 1257

<212> DNA

<213> Megathura crenulata

<400> 54

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 aattggacaa ggcgtcaacaa ctgtcaactgg tggaaagccct cgagtccatg aaagccgacc 120
 attcatctga tgggttccag gcaatcgctt ccttccatgc tcttccatc ctttgcctat 180
 caccagctgc ttcaaaagagg ttgcgtgt gcgtccatgg catggcaacg ttcccacaat 240
 ggcaccgtct gtacacagt caattccaaatg attctctcag aaaacatggt gcagtcgtt 300
 gacttccgtt ctggggatgg acccttccctt gttctgaattt accagagctc ctgaccgtct 360
 caactattca tgaccggag acaggcagag atataccaaatccattt ggttctaaaa 420
 tagagtttga aggagaaaac gtacatacta aaagagatataatcaatggat cgtctcttcc 480
 agggatcaac aaaaacacat cataactggt ttatttgacgca agcactgctt gctcttgaac 540
 aaaccaacta ctgcgacttc gaggttcaatg ttgaaatttgcataatggt gttcataact 600
 ggggtggagg caaggagccc tatggatttgc gcatctgc ttatgtcc tatgtatccac 660
 ttttctacat ccatcaactcc caaaactgtatc gtatgggc tatatggcaa tcgttgcagc 720
 gtttcagagg actttctggta tctgaggctt actgtgtgtt aaatctcatg aaaaactccctc 780
 tgaaggcattt cagtttggta gcaccatata atcttaatgatc acacacgcatttctcaa 840
 agcctgaaga tacattcgac taccaaaatgt ttggatacat atatgacact ctggaaattt 900
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 aatgccttg ggccttgcgttaca agtacgacat aacagaaaacc tttagacaaga 1140
 tgaacccctcg acatgacgaa atcttccaga ttgaagtaac cattacatcc tacgatggaa 1200
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<210> 55

<211> 1254

<212> DNA

<213> Megathura crenulata

<400> 55

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 cagatgggttt tgctgccattt gcatcttccatgc atggatcttgc tgccaaatgtt aatgacgac 180
 acaataaacgaa ggtggcatgc tttatccatg gaatgcctac attcccccac tggcacagac 240
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48

actgggactg gacaaagcca atacataata ttccacatct gttcacagac aaagaataact 360
 acgatgtctg gagaataaa gtaatgccaa atccatttgc ccgagggtat gtcccctcac 420
 acgatacata cacggttaaga gacgtccaag aaggcctgtt ccacctgaca tcaacgggtg 480
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 tccatccctc tgaaccattc ttcatcaagg tgcgttgc agccgtcaac ggaacagttc 1200
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<210> 56

<211> 509

<212> DNA

<213> Megathura crenulata

<400> 56

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 acacacttac cactgcagag gtggacaatc tcaaagatgc catgagagcc gtcatggcag 120
 accacgggtcc aaatggatac caggctatac cagcggttcca tggaaaccca ccaatgttgc 180
 ctatgccaga tggaaagaat tactcgttgc gtacacatgg catggctact ttcccccact 240
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 gccttcctta ctgggacggg acaactgcctt ttagactttt gccaactttt gtcacagatg 360
 aagaggacaa tcccttccat catggtcaca tagactttt gggagtggat acaactcggt 420
 cgccccgaga caagttgttc aatgatccag agcgaggatc agaatcggtc ttctacaggc 480
 aggttcttgc ggccttggag cagacagat 509

<210> 57

<211> 943

<212> DNA

<213> Megathura crenulata

<400> 57

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 gtggcgtttg aaaatgggtt aacaaggcagg aatcctgtat ccaaaactttt tatgaaacca 180
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 ggtcatgata aatacaaaatg gggaaatgtt gtagtacttg gtggatccaa agaaaatgaaa 780
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<210> 58
<211> 1248
<212> DNA
<213> Megathura crenulata

<400> 58

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ggggtttcca gcagattgca gcattccacg gagaaccaa atgtgttcca agccccgaag 180
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<210> 59
<211> 1257
<212> DNA
<213> Megathura crenulata

<400> 59

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aaaaagaggg tatatatgaa aacattgca agttccatgg aaaaaccaggaa ctttgcggaa 180
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gactgtacgt tcttcagggtt gagaatgcgc tcttgcacg agggcttgc gttgtgttgc 300
cttactggaa ctggaccggag aaagctgact ctctgcattt attaatcaat gatgcaactt 360
atttcaattt acgatcccac acctttgatc ctaatcctt cttcaggggaa catattgcct 420
tcgagaatgc tttgttgc agagatccctc agccagaact atggacaat aaggacttct 480
acgagaatgt catgttgcctt ctttgcacg acaacttgc tgacttttag attcagcttgc 540
agctgatataca caacccctt cattcttagac ttggagaaag ggctaaatac tccctttctgt 600
ctcttgatata taccgcattt gatcctgtat ttttccttca ccatgcacac gttgacagaa 660
tctggccat ctggcaggac ttgcagagat atagaagaa accatacaat gaggctgact 720
gcccggatca cgagatgcgt aaaccccttc aaccattaa taacccggaa cttAACAGTG 780
attccatgac gcttaaacac aaccccccac aagacatgt tgattatcaaa aaccgcttca 840
ggtaccaata tgataaacctt caatttaccc acattcgcattt acaaaagctt gaccaaaactt 900
ttcaggcttag aaaaacacac gacagatgtt ttgttgcctt tatttttcac aacatgggaa 960
catctgtgt tttgtatataatggc ttgaaacaaagg aggagaacaa aacttgcac 1020
caaaggccggg ttcccttcacg attctggggg gagaacacaga aatgcattt cactttgacc 1080
gcttgcataaa atttgacata acgtctgcctc tgcataaact ttgttgcctt ttggacggac 1140
atggatttgcg catcaaagtt gacgtcagag ctgtcaatgg atgcatttca gatcaacacaca 1200
tcctcaacgaa accggacttgc ctttttgcgtt ctgttgcacg taagaatata tattatg 1257

50

<210> 60
 <211> 1239
 <212> DNA
 <213> Megathura crenulata

<400> 60
 atggccttc acaacataat cttgtgcgaa aagaagtaag ctctcttaca acactggaga 60
 aacatttttt gaggaaagct ctcaagaaca tgcaagcaga tgattctcca gacggatatc 120
 aagcttattgc ttcttccac gctttgcctc ctctttgtcc aagtcattct gctgcacata 180
 gacacgcttc ttgcctccat ggtatggctc cttccctca gtggcacaga ctctacacag 240
 tttagttcga agatttttga aaacgacatg gttctattgt cggacttcca tattgggatt 300
 ggctgaaacc gcagttcga cttccgtatt tggtagacaca ggagacatac gggcacctgt 360
 tttcacacaa aacctccca aatccgttcc tcaaggcaaa tatagaattt gaggggagagg 420
 gagtaacaac agagagggat gtgtatgtc aacaccttt tgcaaaagga aatctggtt 480
 acaacaactg gttttcaat caggactat atgcactaga acaagaaaat tactgtact 540
 ttgaaataca gttcgaaatt ttgcataatg gaattcattt atgggttgg ggtcaaaaga 600
 cccattcaat aggtcatctt cattacgcat catacgatcc actgttctat atccaccatt 660
 cgcagacaga tcgcatttgg gctatctggc aagcttccca ggagcacaga ggtctttcag 720
 ggaaggaagc acactgcgc ctggagcaaa tggaaagaccc tctcaaacct ttcagctttg 780
 gaagtcccta taatttgaac aaacgcactc aagagtttca caagcctgaa gacacattt 840
 attatcaccg attcgggtat gaggatgatt ccctcgaatt tggtagtgc tctgtttcaa 900
 gtttacataa ctatataaaa caacaacagg aagctgatag agtcttcgca ggattccttc 960
 ttaaaggatt tggacaatca gcatccgtat cgtttgcataatg ctgcagacca gaccagagg 1020
 gccaagaagc tggataacttc tcagttctcg gtggaaagttc agaaatgccc tggcagttg 1080
 acaggctta caagtacgac attacaaaaa ctttgcataatg catgaaactg cgatacgatg 1140
 acacatttac catcaagggtt cacataaagg atatagctgg agttagtgc gacagcgatc 1200
 tgattccaaac tccttctgtt ctccgttgc aaggaaagc 1239

<210> 61
 <211> 1251
 <212> DNA
 <213> Haliotis tuberculata

<400> 61
 atggatcaa tgcgttcac gttggcgta atcggattcg tatggacta tctgaactca 60
 ccgagagaga tctcccccagc ctgaaatctg caatgaggc tctacaagct gacgatgggg 120
 tgaacggta tcaaggcatt gcatcattcc acggcttccc ggcttcttgcatgatgt 180
 agggacatga gattgcgtt tgcgttcacg gaatgccagt attcccacac tggcacaggc 240
 ttacaccct gcaaatggac atggctctgt tatctcacttgc atctgtgtt gctattccat 300
 actggactg gaccaaaacctt atcggccaaac tgcgttgcattt cttcaccaggc cctgaatatt 360
 acgatccttgc gaggatgca gttgtcaata atccatttgc taaaggctac attaaatccg 420
 aggacgcttgc acgggttgc gatcctcagg acatttgtt ccacttgcag gacgaaacgg 480
 gaacatctgt ttgttagat caaactcttt tagccttgc gcaagacatg ttctgtgtt 540
 ttgaggttca atttgggttgc gtcataatg ctatcactt cttgggtgggt ggtcgacaag 600
 ttatgtctt ttcttcttca cactatgtt catatgaccc agccttctt attcatcact 660
 cctttgttgc caaaatatgg gcaatcttgc aagcttgc aagaagaga aagcttccct 720
 atcataaaagc ggattgtgtt ctttacatgc tgaccaaaacc aatgcgacca tttgcacacg 780
 atttcaatca caatgatttcc aaaaaatgc acggacttccc caacactcta ttgtacttcc 840
 aggacctttt ctacacgtt gacaacttag aaattgttgc catgtatgtt aatcgttgg 900
 aagcggaaat caacccggcga aaaagccaaat caagatgtt tgccgggttc cttctacatg 960
 gcattggaaat atcagatgtt gtcacattttt ggatttgc gacagctgac gactgcccacg 1020
 catctggcat gatcttccat ttaggggtt ctttgcataatg gcaactggcc tatgacagaga 1080
 actttaataa ctttgcataatg caatgttgc aggttcacttgc catacaccctt gaaatgttgc 1140
 ttgacacttgc tgcttccat ttcattaaatg tggagggttca tgggttgc aagactgttgc 1200
 tccatcttc agctatccca gcaatcttgc taatctactc agtgggttgc 1251

51

<210> 62
 <211> 1185
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 62
 atcatattgc tggcagtggaa gtcaggaaag acgtgacgac tcttaccgca tctgagatag 60
 agaacctgag gcatgctctg caaagcgtga tggatgtga tggacccaaat ggattccagg 120
 caattgctgc ttatcacggaa agtccttccca tggatgtcat gcntgatggg agagacgtt 180
 catgttgc tcatggaaatg gcatcttcc ctcactggca cagactgttt gtggaaacaga 240
 tggaggatgc actggctgcg catggagctc acattggcat accatactgg gattggacaa 300
 gtgcgttttag tcatctgcct gccttagtga ctgaccacga gcacaatccc ttccaccacg 360
 gacatattgc tcatcgaaat gttggatacat ctgcgtatcc gagagacatg ctgttcaatg 420
 acccccgaaca cgggtcagaa tcattttct atagacaggt tctcttggct ctagaacaga 480
 cagacttctg ccaatttggaa gtcagtttggaaataacaca caatgcaatc cactcttggaa 540
 ctggaggaca tactccatata ggaatgtcat cactggaaata tacagcatat gatccactct 600
 ttatcttccaa ccatttccaaac actgtatcgta tctggccat ctggcaggca ctccagaaat 660
 acagagggtt tcaataacaac gcagcttatttgcgatatcca ggttctgaaa caaccttta 720
 aaccatttcag cgagtcaggaaatccaaacc cagtcaccag agccaaattcttcc 780
 attcatttgc ttatggagaga ctcattatc aatatgacac acttaccccttcc 840
 ctatctcaga acttgatgcc atgcttcaag agagaaagaa ggaagagaga acatttgc 900
 cttccctgtt gcacggattt ggcgccttgc ctgtatgttc gtttgcaccccttcc 960
 atggtcatttgc ttatggagaga ctcattatc aatatgacac acttaccccttcc 1020
 gtccttgc aagatttgc cgttacgata tcacaaaggt tctcaagcag atgaatcttcc 1080
 actatgatttgc tgagttccat ttggatgttgc agatttgc cacagatggaa acagaactgc 1140
 catcgatcg tatcaagagc cctaccatttgc aacaccatgg aggag 1185

<210> 63
 <211> 422
 <212> PRT
 <213> *Haliotis tuberculata*

<220>
 <221> SIGNAL
 <222> (1)...(15)

<400> 63
 Leu Val Gln Phe Leu Leu Val Ala Leu Val Val Gly Ala Gly Ala Asp
 1 5 10 15
 Asn Val Val Arg Lys Asp Val Ser His Leu Thr Asp Asp Glu Val Gln
 20 25 30
 Ala Leu His Gly Ala Leu His Asp Val Thr Ala Ser Thr Gly Pro Leu
 35 40 45
 Ser Phe Glu Asp Ile Thr Ser Tyr His Ala Ala Pro Ala Ser Cys Asp
 50 55 60
 Tyr Lys Gly Arg Lys Ile Ala Cys Cys Val His Gly Met Pro Ser Phe
 65 70 75 80
 Pro Phe Trp His Arg Ala Tyr Val Val Gln Ala Glu Arg Ala Leu Leu
 85 90 95
 Ser Lys Arg Lys Thr Val Gly Met Pro Tyr Trp Asp Trp Thr Gln Thr
 100 105 110

Leu Thr His Leu Pro Ser Leu Val Thr Glu Pro Ile Tyr Ile Asp Ser
 115 120 125
 Lys Gly Gly Lys Ala Gln Thr Asn Tyr Trp Tyr Arg Gly Glu Ile Ala
 130 135 140
 Phe Ile Asn Lys Lys Thr Ala Arg Ala Val Asp Asp Arg Leu Phe Glu
 145 150 155 160
 Lys Val Glu Pro Gly His Tyr Thr His Leu Met Glu Thr Val Leu Asp
 165 170 175
 Ala Leu Glu Gln Asp Glu Phe Cys Lys Phe Glu Ile Gln Phe Glu Leu
 180 185 190
 Ala His Asn Ala Ile His Tyr Leu Val Gly Gly Lys Phe Glu Tyr Ser
 195 200 205
 Met Ser Asn Leu Glu Tyr Thr Ser Tyr Asp Pro Ile Phe Phe Leu His
 210 215 220
 His Ser Asn Val Asp Arg Leu Phe Ala Ile Trp Gln Arg Leu Gln Glu
 225 230 235 240
 Leu Arg Gly Lys Asn Pro Asn Ala Met Asp Cys Ala His Glu Leu Ala
 245 250 255
 His Gln Gln Leu Gln Pro Phe Asn Arg Asp Ser Asn Pro Val Gln Leu
 260 265 270
 Thr Lys Asp His Ser Thr Pro Ala Asp Leu Phe Asp Tyr Lys Gln Leu
 275 280 285
 Gly Tyr Ser Tyr Asp Ser Leu Asn Leu Asn Gly Met Thr Pro Glu Gln
 290 295 300
 Leu Lys Thr Glu Leu Asp Glu Arg His Ser Lys Glu Arg Ala Phe Ala
 305 310 315 320
 Ser Phe Arg Leu Ser Gly Phe Gly Ser Ala Asn Val Val Tyr
 325 330 335
 Ala Cys Val Pro Asp Asp Pro Arg Ser Asp Asp Tyr Cys Glu Lys
 340 345 350
 Ala Gly Asp Phe Phe Ile Leu Gly Gln Ser Glu Met Pro Trp Arg
 355 360 365
 Phe Tyr Arg Pro Phe Phe Tyr Asp Val Thr Glu Ala Val His His Leu
 370 375 380
 Gly Val Pro Leu Ser Gly His Tyr Tyr Val Lys Thr Glu Leu Phe Ser
 385 390 395 400
 Val Asn Gly Thr Ala Leu Ser Pro Asp Leu Leu Pro Gln Pro Thr Val
 405 410 415

Ala Tyr Arg Pro Gly Lys
420

<210> 64
<211> 511
<212> PRT
<213> *Haliotis tuberculata*

<400> 64
Val His Arg Gly Gly Asn His Glu Asp Glu His His Asp Asp Arg Leu
1 5 10 15

Ala Asp Val Leu Ile Arg Lys Glu Val Asp Phe Leu Ser Leu Gln Glu
20 25 30

Ala Asn Ala Ile Lys Asp Ala Leu Tyr Lys Leu Gln Asn Asp Asp Ser
35 40 45

Lys Gly Gly Phe Glu Ala Ile Ala Gly Tyr His Gly Tyr Pro Asn Met
50 55 60

Cys Pro Glu Arg Gly Thr Asp Lys Tyr Pro Cys Cys Val His Gly Met
65 70 75 80

Pro Val Phe Pro His Trp His Arg Leu His Thr Ile Gln Met Glu Arg
85 90 95

Ala Leu Lys Asn His Gly Ser Pro Met Gly Ile Pro Tyr Trp Asp Trp
100 105 110

Thr Lys Lys Met Ser Ser Leu Pro Ser Phe Phe Gly Asp Ser Ser Asn
115 120 125

Asn Asn Pro Phe Tyr Lys Tyr Ile Arg Gly Val Gln His Glu Thr
130 135 140

Thr Arg Asp Val Asn Gln Arg Leu Phe Asn Gln Thr Lys Phe Gly Glu
145 150 155 160

Phe Asp Tyr Leu Tyr Tyr Leu Thr Leu Gln Val Leu Glu Glu Asn Ser
165 170 175

Tyr Cys Asp Phe Glu Val Gln Tyr Glu Ile Leu His Asn Ala Val His
180 185 190

Ser Trp Leu Gly Gly Thr Gly Gln Tyr Ser Met Ser Thr Leu Glu His
195 200 205

Ser Ala Phe Asp Pro Val Phe Met Ile His His Ser Ser Leu Asp Arg
210 215 220

Ile Trp Ile Leu Trp Gln Lys Leu Gln Lys Ile Arg Met Lys Pro Tyr
225 230 235 240

54

Tyr Ala Leu Asp Cys Ala Gly Asp Arg Leu Met Lys Asp Pro Leu His
 245 250 255

Pro Phe Asn Tyr Glu Thr Val Asn Glu Asp Glu Phe Thr Arg Ile Asn
 260 265 270

Ser Phe Pro Ser Ile Leu Phe Asp His Tyr Arg Phe Asn Tyr Glu Tyr
 275 280 285

Asp Asn Met Arg Ile Arg Gly Gln Asp Ile His Glu Leu Glu Glu Val
 290 295 300

Ile Gln Glu Leu Arg Asn Lys Asp Arg Ile Phe Ala Gly Phe Val Leu
 305 310 315 320

Ser Gly Leu Arg Ile Ser Ala Thr Val Lys Val Phe Ile His Ser Lys
 325 330 335

Asn Asp Thr Ser His Glu Glu Tyr Ala Gly Glu Phe Ala Val Leu Gly
 340 345 350

Gly Glu Lys Glu Met Pro Trp Ala Tyr Glu Arg Met Leu Lys Leu Asp
 355 360 365

Ile Ser Asp Ala Val His Lys Leu His Val Lys Asp Glu Asp Ile Arg
 370 375 380

Phe Arg Val Val Val Thr Ala Tyr Asn Gly Asp Val Val Thr Thr Arg
 385 390 395 400

Leu Ser Gln Pro Phe Ile Val His Arg Pro Ala His Val Ala His Asp
 405 410 415

Ile Leu Val Ile Pro Val Gly Ala Gly His Asp Leu Pro Pro Lys Val
 420 425 430

Val Val Lys Ser Gly Thr Lys Val Glu Phe Thr Pro Ile Asp Ser Ser
 435 440 445

Val Asn Lys Ala Met Val Glu Leu Gly Ser Tyr Thr Ala Met Ala Lys
 450 455 460

Cys Ile Val Pro Pro Phe Ser Tyr His Gly Phe Glu Leu Asp Lys Val
 465 470 475 480

Tyr Ser Val Asp His Gly Asp Tyr Tyr Ile Ala Ala Gly Thr His Ala
 485 490 495

Leu Cys Glu Gln Asn Leu Arg Leu His Ile His Val Glu His Glu
 500 505 510

<210> 65
 <211> 197
 <212> PRT
 <213> *Haliothis tuberculata*

<400> 65 55
 Gly Leu Pro Tyr Trp Asp Trp Thr Gln His Leu Thr Gln Leu Pro Asp
 1 5 10 15
 Leu Val Ser Asp Pro Leu Phe Val Asp Pro Glu Gly Gly Lys Ala His
 20 25 30
 Asp Asn Ala Trp Tyr Arg Gly Asn Ile Lys Phe Glu Asn Lys Lys Thr
 35 40 45
 Ala Arg Ala Val Asp Asp Arg Leu Phe Glu Lys Val Gly Pro Gly Glu
 50 55 60
 Asn Thr Arg Leu Phe Glu Gly Ile Leu Asp Ala Leu Glu Gln Asp Glu
 65 70 75 80
 Phe Cys Asn Phe Glu Ile Gln Phe Glu Leu Ala His Asn Ala Ile His
 85 90 95
 Tyr Leu Val Gly Gly Arg His Thr Tyr Ser Met Ser His Leu Glu Tyr
 100 105 110
 Thr Ser Tyr Asp Pro Leu Phe Phe Leu His His Ser Asn Pro Asp Arg
 115 120 125
 Ile Phe Ala Ile Trp Glu Arg Leu Gln Val Leu Arg Gly Lys Asp Pro
 130 135 140
 Asn Thr Ala Asp Cys Ala His Asn Leu Ile His Glu Pro Met Glu Pro
 145 150 155 160
 Phe Arg Arg His Glu Pro Met Glu Pro Phe Arg Arg Asp Ser Asn Pro
 165 170 175
 Leu Asp Leu Thr Arg Glu Asn Ser Lys Pro Ile Asp Ser Phe Asp Tyr
 180 185 190
 Ala His Leu Gly Tyr
 195

<210> 66
 <211> 415
 <212> PRT
 <213> Haliotis tuberculata

<400> 66
 Val Thr Glu Ala Pro Ala Pro Ser Ser Asp Ala His Leu Ala Val Arg
 1 5 10 15
 Lys Asp Ile Asn His Leu Thr Arg Glu Glu Val Tyr Glu Leu Arg Arg
 20 25 30
 Ala Met Glu Arg Phe Gln Ala Asp Thr Ser Val Asp Gly Tyr Gln Ala
 35 40 45

56

Thr Val Glu Tyr His Gly Leu Pro Ala Arg Cys Pro Phe Pro Glu Ala
 50 55 60

Thr Asn Arg Phe Ala Cys Cys Ile His Gly Met Ala Thr Phe Pro His
 65 70 75 80

Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Ile Arg Arg
 85 90 95

Gly Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Gln Pro Met Ala
 100 105 110

His Leu Pro Gly Leu Ala Asp Asn Ala Thr Tyr Arg Asp Pro Ile Ser
 115 120 125

Gly Asp Ser Arg His Asn Pro Phe His Asp Val Glu Val Ala Phe Glu
 130 135 140

Asn Gly Arg Thr Glu Arg His Pro Asp Ser Arg Leu Phe Glu Gln Pro
 145 150 155 160

Leu Phe Gly Lys His Thr Arg Leu Phe Asp Ser Ile Val Tyr Ala Phe
 165 170 175

Glu Gln Glu Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Met Thr His
 180 185 190

Asn Asn Ile His Ala Trp Ile Gly Gly Glu Lys Tyr Ser Met Ser
 195 200 205

Ser Leu His Tyr Thr Ala Phe Asp Pro Ile Phe Tyr Leu Arg His Ser
 210 215 220

Asn Thr Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg
 225 230 235 240

Asn Arg Pro Tyr Lys Ala His Cys Ala Trp Ser Glu Glu Arg Gln Pro
 245 250 255

Leu Lys Pro Phe Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr
 260 265 270

Tyr Glu Asn Ser Val Pro Thr Asn Val Tyr Asp Tyr Glu Gly Val Leu
 275 280 285

Gly Tyr Thr Tyr Asp Asp Leu Asn Phe Gly Gly Met Asp Leu Gly Gln
 290 295 300

Leu Glu Glu Tyr Ile Gln Arg Gln Arg Asp Arg Thr Phe Ala
 305 310 315 320

Gly Phe Phe Leu Ser His Ile Gly Thr Ser Ala Asn Val Glu Ile Ile
 325 330 335

Ile Asp His Gly Thr Leu His Thr Ser Val Gly Thr Phe Ala Val Leu
 340 345 350

Gly Gly Glu Lys Glu Met Lys Trp Gly Phe Asp Arg Leu Tyr Lys Tyr 57
 355 360 365
 Glu Ile Thr Asp Glu Leu Arg Gln Leu Asn Leu Arg Ala Asp Asp Val
 370 375 380
 Phe Ser Ile Ser Val Lys Val Thr Asp Val Asp Gly Ser Glu Leu Ser
 385 390 395 400
 Ser Glu Leu Ile Pro Ser Ala Ala Ile Ile Phe Glu Arg Ser His
 405 410 415
 <210> 67
 <211> 414
 <212> PRT
 <213> *Haliothis tuberculata*
 <400> 67
 Gly His His Gln Ala Asp Glu Tyr Asp Glu Val Val Thr Ala Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Leu Lys Asp Leu Ser Lys Gly Glu Val Glu Ser
 20 25 30
 Leu Arg Ser Ala Phe Leu Gln Leu Gln Asn Asp Gly Val Tyr Glu Asn
 35 40 45
 Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Asp Asp Asn Gly Arg
 50 55 60
 Lys Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
 65 70 75 80
 Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95
 Ala Val Ser Val Pro Tyr Trp Asp Trp Thr Glu Thr Phe Thr Glu Leu
 100 105 110
 Pro Ser Leu Ile Ala Glu Ala Thr Tyr Phe Asn Ser Arg Gln Gln Thr
 115 120 125
 Phe Asp Pro Asn Pro Phe Phe Arg Gly Lys Ile Ser Phe Glu Asn Ala
 130 135 140
 Val Thr Thr Arg Asp Pro Gln Pro Glu Leu Tyr Val Asn Arg Tyr Tyr
 145 150 155 160
 Tyr Gln Asn Val Met Leu Val Phe Glu Gln Asp Asn Tyr Cys Asp Phe
 165 170 175
 Glu Ile Gln Phe Glu Met Val His Asn Val Leu His Ala Trp Leu Gly
 180 185 190
 Gly Arg Ala Thr Tyr Ser Ile Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205

Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220

Trp Gln Glu Leu Gln Arg Tyr Arg Lys Lys Pro Tyr Asn Glu Ala Asp
 225 230 235 240

Cys Ala Ile Asn Leu Met Arg Lys Pro Leu His Pro Phe Asp Asn Ser
 245 250 255

Asp Leu Asn His Asp Pro Val Thr Phe Lys Tyr Ser Lys Pro Thr Asp
 260 265 270

Gly Phe Asp Tyr Gln Asn Asn Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285

Phe Asn His Phe Ser Ile Pro Arg Leu Glu Glu Ile Arg Ile Arg
 290 295 300

Gln Arg Gln Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320

Thr Ser Ala Thr Val Glu Ile Phe Val Cys Val Pro Thr Thr Ser Gly
 325 330 335

Glu Gln Asn Cys Glu Asn Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 340 345 350

Glu Thr Glu Met Ala Phe His Phe Asp Arg Leu Tyr Arg Phe Asp Ile
 355 360 365

Ser Glu Thr Leu Arg Asp Leu Gly Ile Gln Leu Asp Ser His Asp Phe
 370 375 380

Asp Leu Ser Ile Lys Ile Gln Gly Val Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400

His Ile Leu Pro Glu Pro Ser Leu Ile Phe Val Pro Gly Ser
 405 410

<210> 68
 <211> 419
 <212> PRT
 <213> *Haliotis tuberculata*

<400> 68
 Ser Ser Phe Leu Arg Pro Asp Gly His Ser Asp Asp Ile Leu Val Arg
 1 5 10 15

Lys Glu Val Asn Ser Leu Thr Thr Arg Glu Thr Ala Ser Leu Ile His
 20 25 30

Ala Leu Lys Ser Met Gln Glu Asp His Ser Pro Asp Gly Phe Gln Ala
 35 40 45

59

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala
 50 55 60

Ala His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln
 65 70 75 80

Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ala Leu Arg Arg His
 85 90 95

Gly Ala Thr Val Gly Val Pro Tyr Trp Asp Trp Leu Arg Pro Gln Ser
 100 105 110

His Leu Pro Glu Leu Val Thr Met Glu Thr Tyr His Asp Ile Trp Ser
 115 120 125

Asn Arg Asp Phe Pro Asn Pro Phe Tyr Gln Ala Asn Ile Glu Phe Glu
 130 135 140

Gly Glu Asn Ile Thr Thr Glu Arg Glu Val Ile Ala Asp Lys Leu Phe
 145 150 155 160

Val Lys Gly Gly His Val Phe Asp Lys Leu Val Leu Gln Thr Ser His
 165 170 175

Pro Ser Ala Glu Gln Glu Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu
 180 185 190

Ile Leu His Asn Gly Val His Thr Trp Val Gly Gly Ser Arg Thr Tyr
 195 200 205

Ser Ile Gly His Leu His Tyr Ala Phe Tyr Asp Pro Leu Phe Tyr Leu
 210 215 220

His His Phe Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu Gln
 225 230 235 240

Glu Gln Arg Gly Leu Ser Gly Asp Glu Ala His Cys Ala Leu Glu Gln
 245 250 255

Met Arg Glu Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Trp
 260 265 270

Asn Gln Leu Thr Gln Asp Phe Ser Arg Pro Glu Asp Thr Phe Asp Tyr
 275 280 285

Arg Lys Phe Gly Tyr Glu Tyr Asp Asn Leu Glu Phe Leu Gly Met Ser
 290 295 300

Val Ala Glu Leu Asp Gln Tyr Ile Ile Glu His Gln Glu Asn Asp Arg
 305 310 315 320

Val Phe Ala Gly Phe Leu Leu Ser Gly Phe Gly Gly Ser Ala Ser Val
 325 330 335

Asn Phe Gln Val Cys Arg Ala Asp Ser Thr Cys Gln Asp Ala Gly Tyr
 340 345 350

Phe Thr Val Leu Gly Gly Ser Ala Glu Met Ala Trp Ala Phe Asp Arg 60
 355 360 365
 Leu Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Glu Lys Met His Leu Arg
 370 375 380
 Tyr Asp Asp Asp Phe Thr Ile Ser Val Ser Leu Thr Ala Asn Asn Gly
 385 390 395 400
 Thr Val Leu Ser Ser Leu Ile Pro Thr Pro Ser Val Ile Phe Gln
 405 410 415
 Arg Gly His

<210> 69
 <211> 378
 <212> PRT
 <213> Megathura crenulata

<400> 69
 Arg Tyr Gln Ala Thr Ala Glu Tyr His Gly Leu Pro Ala Arg Cys Pro
 1 5 10 15
 Arg Pro Asp Ala Lys Asp Arg Tyr Ala Cys Cys Val His Gly Met Pro
 20 25 30
 Ile Phe Pro His Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala
 35 40 45
 Leu Val Gly Arg Gly Ala Thr Ile Gly Ile Pro Tyr Trp Asp Trp Thr
 50 55 60
 Glu Pro Met Thr His Ile Pro Gly Leu Ala Gly Asn Lys Thr Tyr Val
 65 70 75 80
 Asp Ser His Gly Ala Ser His Thr Asn Pro Phe His Ser Ser Val Ile
 85 90 95
 Ala Phe Glu Glu Asn Ala Pro His Thr Lys Arg Gln Ile Asp Gln Arg
 100 105 110
 Leu Phe Lys Pro Ala Thr Phe Gly His His Thr Asp Leu Phe Asn Gln
 115 120 125
 Ile Leu Tyr Ala Phe Glu Gln Glu Asp Tyr Cys Asp Phe Glu Val Gln
 130 135 140
 Phe Glu Ile Thr His Asn Thr Ile His Ala Trp Thr Gly Gly Ser Glu
 145 150 155 160
 His Phe Ser Met Ser Ser Leu His Tyr Thr Ala Phe Asp Pro Leu Phe
 165 170 175
 Tyr Phe His His Ser Asn Val Asp Arg Leu Trp Ala Val Trp Gln Ala
 180 185 190

Leu Gln Met Arg Arg His Lys Pro Tyr Arg Ala His Cys Ala Ile Ser
 195 200 205

Leu Glu His Met His Leu Lys Pro Phe Ala Phe Ser Ser Pro Leu Asn
 210 215 220

Asn Asn Glu Lys Thr His Ala Asn Ala Met Pro Asn Lys Ile Tyr Asp
 225 230 235 240

Tyr Glu Asn Val Leu His Tyr Thr Tyr Glu Asp Leu Thr Phe Gly Gly
 245 250 255

Ile Ser Leu Glu Asn Ile Glu Lys Met Ile His Glu Asn Gln Gln Glu
 260 265 270

Asp Arg Ile Tyr Ala Gly Phe Leu Leu Ala Gly Ile Arg Thr Ser Ala
 275 280 285

Asn Val Asp Ile Phe Ile Lys Thr Thr Asp Ser Val Gln His Lys Ala
 290 295 300

Gly Thr Phe Ala Val Leu Gly Gly Ser Lys Glu Met Lys Trp Gly Phe
 305 310 315 320

Asp Arg Val Phe Lys Phe Asp Ile Thr His Val Leu Lys Asp Leu Asp
 325 330 335

Leu Thr Ala Asp Gly Asp Phe Glu Val Thr Val Asp Ile Thr Glu Val
 340 345 350

Asp Gly Thr Lys Leu Ala Ser Ser Leu Ile Pro His Ala Ser Val Ile
 355 360 365

Arg Glu His Ala Arg Gly Lys Leu Asn Arg
 370 375

<210> 70

<211> 419

<212> PRT

<213> Megathura crenulata

<400> 70

Asp Ser Ala His Thr Asp Asp Gly His Thr Glu Pro Val Met Ile Arg
 1 5 10 15

Lys Asp Ile Thr Gln Leu Asp Lys Arg Gln Gln Leu Ser Leu Val Lys
 20 25 30

Ala Leu Glu Ser Met Lys Ala Asp His Ser Ser Asp Gly Phe Gln Ala
 35 40 45

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ala Ala
 50 55 60

Ser Lys Arg Phe Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln 62
 65 70 75 80
 Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ser Leu Arg Lys His
 85 90 95
 Gly Ala Val Val Gly Leu Pro Tyr Trp Asp Trp Thr Leu Pro Arg Ser
 100 105 110
 Glu Leu Pro Glu Leu Leu Thr Val Ser Thr Ile His Asp Pro Glu Thr
 115 120 125
 Gly Arg Asp Ile Pro Asn Pro Phe Ile Gly Ser Lys Ile Glu Phe Glu
 130 135 140
 Gly Glu Asn Val His Thr Lys Arg Asp Ile Asn Arg Asp Arg Leu Phe
 145 150 155 160
 Gln Gly Ser Thr Lys Thr His His Asn Trp Phe Ile Glu Gln Ala Leu
 165 170 175
 Leu Ala Leu Glu Gln Thr Asn Tyr Cys Asp Phe Glu Val Gln Phe Glu
 180 185 190
 Ile Met His Asn Gly Val His Thr Trp Val Gly Gly Lys Glu Pro Tyr
 195 200 205
 Gly Ile Gly His Leu His Tyr Ala Ser Tyr Asp Pro Leu Phe Tyr Ile
 210 215 220
 His His Ser Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Ser Leu Gln
 225 230 235 240
 Arg Phe Arg Gly Leu Ser Gly Ser Glu Ala Asn Cys Ala Val Asn Leu
 245 250 255
 Met Lys Thr Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Leu
 260 265 270
 Asn Asp His Thr His Asp Phe Ser Lys Pro Glu Asp Thr Phe Asp Tyr
 275 280 285
 Gln Lys Phe Gly Tyr Ile Tyr Asp Thr Leu Glu Phe Ala Gly Trp Ser
 290 295 300
 Ile Arg Gly Ile Asp His Ile Val Arg Asn Arg Gln Glu His Ser Arg
 305 310 315 320
 Val Phe Ala Gly Phe Leu Leu Glu Gly Phe Gly Thr Ser Ala Thr Val
 325 330 335
 Asp Phe Gln Val Cys Arg Thr Ala Gly Asp Cys Glu Asp Ala Gly Tyr
 340 345 350
 Phe Thr Val Leu Gly Gly Glu Lys Glu Met Pro Trp Ala Phe Asp Arg
 355 360 365

63

Leu Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Asp Lys Met Asn Leu Arg
 370 375 380

His Asp Glu Ile Phe Gln Ile Glu Val Thr Ile Thr Ser Tyr Asp Gly
 385 390 395 400

Thr Val Leu Asp Ser Gly Leu Ile Pro Thr Pro Ser Ile Ile Tyr Asp
 405 410 415

Pro Ala His

<210> 71

<211> 418

<212> PRT

<213> Megathura crenulata

<400> 71

His Asp Ile Ser Ser His His Leu Ser Leu Asn Lys Val Arg His Asp
 1 5 10 15

Leu Ser Thr Leu Ser Glu Arg Asp Ile Gly Ser Leu Lys Tyr Ala Leu
 20 25 30

Ser Ser Leu Gln Ala Asp Thr Ser Ala Asp Gly Phe Ala Ala Ile Ala
 35 40 45

Ser Phe His Gly Leu Pro Ala Lys Cys Asn Asp Ser His Asn Asn Glu
 50 55 60

Val Ala Cys Cys Ile His Gly Met Pro Thr Phe Pro His Trp His Arg
 65 70 75 80

Leu Tyr Thr Leu Gln Phe Glu Gln Ala Leu Arg Arg His Gly Ser Ser
 85 90 95

Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Pro Ile His Asn Ile Pro
 100 105 110

His Leu Phe Thr Asp Lys Glu Tyr Tyr Asp Val Trp Arg Asn Lys Val
 115 120 125

Met Pro Asn Pro Phe Ala Arg Gly Tyr Val Pro Ser His Asp Thr Tyr
 130 135 140

Thr Val Arg Asp Val Gln Glu Gly Leu Phe His Leu Thr Ser Thr Gly
 145 150 155 160

Glu His Ser Ala Leu Leu Asn Gln Ala Leu Leu Ala Leu Glu Gln His
 165 170 175

Asp Tyr Cys Asp Phe Ala Val Gln Phe Glu Val Met His Asn Thr Ile
 180 185 190

His Tyr Leu Val Gly Gly Pro Gln Val Tyr Ser Leu Ser Ser Leu His
 195 200 205

Tyr Ala Ser Tyr Asp Pro Ile Phe Phe Ile His His Ser Phe Val Asp
 210 215 220
 Lys Val Trp Ala Val Trp Gln Ala Leu Gln Glu Lys Arg Gly Leu Pro
 225 230 235 240
 Ser Asp Arg Ala Asp Cys Ala Val Ser Leu Met Thr Gln Asn Met Arg
 245 250 255
 Pro Phe His Tyr Glu Ile Asn His Asn Gln Phe Thr Lys Lys His Ala
 260 265 270
 Val Pro Asn Asp Val Phe Lys Tyr Glu Leu Leu Gly Tyr Arg Tyr Asp
 275 280 285
 Asn Leu Glu Ile Gly Gly Met Asn Leu His Glu Ile Glu Lys Glu Ile
 290 295 300
 Lys Asp Lys Gln His His Val Arg Val Phe Ala Gly Phe Leu Leu His
 305 310 315 320
 Gly Ile Arg Thr Ser Ala Asp Val Gln Phe Gln Ile Cys Lys Thr Ser
 325 330 335
 Glu Asp Cys His His Gly Gly Gln Ile Phe Val Leu Gly Gly Thr Lys
 340 345 350
 Glu Met Ala Trp Ala Tyr Asn Arg Leu Phe Lys Tyr Asp Ile Thr His
 355 360 365
 Ala Leu His Asp Ala His Ile Thr Pro Glu Asp Val Phe His Pro Ser
 370 375 380
 Glu Pro Phe Phe Ile Lys Val Ser Val Thr Ala Val Asn Gly Thr Val
 385 390 395 400
 Leu Pro Ala Ser Ile Leu His Ala Pro Thr Ile Ile Tyr Glu Pro Gly
 405 410 415
 Leu Gly

<210> 72
 <211> 241
 <212> PRT
 <213> Megathura crenulata

<400> 72
 Asp His His Glu Asp His His Ser Ser Ser Met Ala Gly His Gly Val
 1 5 10 15
 Arg Lys Glu Ile Asn Thr Leu Thr Thr Ala Glu Val Asp Asn Leu Lys
 20 25 30

65

Asp Ala Met Arg Ala Val Met Ala Asp His Gly Pro Asn Gly Tyr Gln
 35 40 45

Ala Ile Ala Ala Phe His Gly Asn Pro Pro Met Cys Pro Met Pro Asp
 50 55 60

Gly Lys Asn Tyr Ser Cys Cys Thr His Gly Met Ala Thr Phe Pro His
 65 70 75 80

Trp His Arg Leu Tyr Thr Lys Gln Met Glu Asp Ala Leu Thr Ala His
 85 90 95

Gly Ala Arg Val Gly Leu Pro Tyr Trp Asp Gly Thr Thr Ala Phe Thr
 100 105 110

Ala Leu Pro Thr Phe Val Thr Asp Glu Glu Asp Asn Pro Phe His His
 115 120 125

Gly His Ile Asp Tyr Leu Gly Val Asp Thr Thr Arg Ser Pro Arg Asp
 130 135 140

Lys Leu Phe Asn Asp Pro Glu Arg Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160

Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Phe Cys Gln Phe Glu Val
 165 170 175

Gln Phe Glu Ile Thr His Asn Ala Ile His Ser Trp Thr Gly Gly Leu
 180 185 190

Thr Pro Tyr Gly Met Ser Thr Leu Glu Tyr Thr Thr Tyr Asp Pro Leu
 195 200 205

Phe Trp Leu His His Ala Asn Thr Asp Arg Ile Trp Ala Ile Trp Gln
 210 215 220

Ala Leu Gln Glu Tyr Arg Gly Leu Pro Tyr Asp His Ala Asn Cys Glu
 225 230 235 240

Ile

<210> 73

<211> 98

<212> PRT

<213> Megathura crenulata

<400> 73

Lys His His Glu Lys His His Glu Asp His His Glu Asp Ile Leu Val
 1 5 10 15

Arg Lys Asn Ile His Ser Leu Ser His His Glu Ala Glu Glu Leu Arg
 20 25 30

Asp Ala Leu Tyr Lys Leu Gln Asn Asp Glu Ser His Gly Gly Tyr Glu
 35 40 45

66

His Ile Ala Gly Phe His Gly Tyr Pro Asn Leu Cys Pro Glu Lys Gly
 50 55 60
 Asp Glu Lys Tyr Pro Cys Cys Val His Gly Met Ser Ile Phe Pro His
 65 70 75 80
 Trp His Arg Leu His Thr Ile Gln Leu Glu Arg Ala Leu Lys Lys His
 85 90 95
 Gly Ser

<210> 74
 <211> 314
 <212> PRT
 <213> Megathura crenulata

<400> 74
 Gly Leu Pro Tyr Trp Asp Trp Thr Met Pro Met Ser His Leu Pro Glu
 1 5 10 15
 Leu Ala Thr Ser Glu Thr Tyr Leu Asp Pro Val Thr Gly Glu Thr Lys
 20 25 30
 Asn Asn Pro Phe His His Ala Gln Val Ala Phe Glu Asn Gly Val Thr
 35 40 45
 Ser Arg Asn Pro Asp Ala Lys Leu Phe Met Lys Pro Thr Tyr Gly Asp
 50 55 60
 His Thr Tyr Leu Phe Asp Ser Met Ile Tyr Ala Phe Glu Gln Glu Asp
 65 70 75 80
 Phe Cys Asp Phe Glu Val Gln Tyr Glu Leu Thr His Asn Ala Ile His
 85 90 95
 Ala Trp Val Gly Gly Ser Glu Lys Tyr Ser Met Ser Ser Leu His Tyr
 100 105 110
 Thr Ala Phe Asp Pro Ile Phe Tyr Leu His His Ser Asn Val Asp Arg
 115 120 125
 Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg Gly Lys Ser Tyr
 130 135 140
 Lys Ala His Cys Ala Ser Ser Gln Glu Arg Glu Pro Leu Lys Pro Phe
 145 150 155 160
 Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr Tyr His Asn Ser
 165 170 175
 Val Pro Thr Asn Val Tyr Asp Tyr Val Gly Val Leu His Tyr Arg Tyr
 180 185 190

67

Asp Asp Leu Gln Phe Gly Gly Met Thr Met Ser Glu Leu Glu Glu Tyr
 195 200 205

Ile His Lys Gln Thr Gln His Asp Arg Thr Phe Ala Gly Phe Phe Leu
 210 215 220

Ser Tyr Ile Gly Thr Ser Ala Ser Val Asp Ile Phe Ile Asn Arg Glu
 225 230 235 240

Gly His Asp Lys Tyr Lys Val Gly Ser Phe Val Val Leu Gly Gly Ser
 245 250 255

Lys Glu Met Lys Trp Gly Phe Asp Arg Met Tyr Lys Tyr Glu Ile Thr
 260 265 270

Glu Ala Leu Lys Thr Leu Asn Val Ala Val Asp Asp Gly Phe Ser Ile
 275 280 285

Thr Val Glu Ile Thr Asp Val Asp Gly Ser Pro Pro Ser Ala Asp Leu
 290 295 300

Ile Pro Pro Pro Ala Ile Ile Phe Glu Arg
 305 310

<210> 75
 <211> 416
 <212> PRT
 <213> Megathura crenulata

<400> 75

Ala Asp Ala Lys Asp Phe Gly His Ser Arg Lys Ile Arg Lys Ala Val
 1 5 10 15

Asp Ser Leu Thr Val Glu Glu Gln Thr Ser Leu Arg Arg Ala Met Ala
 20 25 30

Asp Leu Gln Asp Asp Lys Thr Ser Gly Gly Phe Gln Gln Ile Ala Ala
 35 40 45

Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Glu Lys Lys
 50 55 60

Phe Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
 65 70 75 80

Leu Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Phe Thr
 85 90 95

Gly Gly Leu Pro Tyr Trp Asp Trp Thr Arg Ser Met Ser Ala Leu Pro
 100 105 110

His Phe Val Ala Asp Pro Thr Tyr Asn Asp Ala Ile Ser Ser Gln Glu
 115 120 125

Glu Asp Asn Pro Trp His His Gly His Ile Asp Ser Val Gly His Asp
 130 135 140

Thr Thr Arg Asp Val Arg Asp Asp Leu Tyr Gln Ser Pro Gly Phe Gly
 145 150 155 160
 His Tyr Thr Asp Ile Ala Lys Gln Val Leu Leu Ala Phe Glu Gln Asp
 165 170 175
 Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn Phe Ile
 180 185 190
 His Ala Leu Val Gly Gly Asn Glu Pro Tyr Ser Met Ser Ser Leu Arg
 195 200 205
 Tyr Thr Thr Tyr Asp Pro Ile Phe Phe Leu His Arg Ser Asn Thr Asp
 210 215 220
 Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro
 225 230 235 240
 Tyr Asn Thr Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro Leu Gln
 245 250 255
 Pro Phe Gly Leu Asp Ser Val Ile Asn Pro Asp Asp Glu Thr Arg Glu
 260 265 270
 His Ser Val Pro Phe Arg Val Phe Asp Tyr Lys Asn Asn Phe Asp Tyr
 275 280 285
 Glu Tyr Glu Ser Leu Ala Phe Asn Gly Leu Ser Ile Ala Gln Leu Asp
 290 295 300
 Arg Glu Leu Gln Arg Arg Lys Ser His Asp Arg Val Phe Ala Gly Phe
 305 310 315 320
 Leu Leu His Glu Ile Gly Gln Ser Ala Leu Val Lys Phe Tyr Val Cys
 325 330 335
 Lys His Asn Val Ser Asp Cys Asp His Tyr Ala Gly Glu Phe Tyr Ile
 340 345 350
 Leu Gly Asp Glu Ala Glu Met Pro Trp Arg Tyr Asp Arg Val Tyr Lys
 355 360 365
 Tyr Glu Ile Thr Gln Gln Leu His Asp Leu Asp Leu His Val Gly Asp
 370 375 380
 Asn Phe Phe Leu Lys Tyr Glu Ala Phe Asp Leu Asn Gly Gly Ser Leu
 385 390 395 400
 Gly Gly Ser Ile Phe Ser Gln Pro Ser Val Ile Phe Glu Pro Ala Ala
 405 410 415

<210> 76
 <211> 419
 <212> PRT

69

<213> Megathura crenulata

<400> 76

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Ser | His | Gln | Ala | Asp | Glu | Tyr | Arg | Glu | Ala | Val | Thr | Ser | Ala | Ser |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| His | Ile | Arg | Lys | Asn | Ile | Arg | Asp | Leu | Ser | Glu | Gly | Glu | Ile | Glu | Ser |
| | | | | | 20 | | | 25 | | | | | 30 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Arg | Ser | Ala | Phe | Leu | Gln | Ile | Gln | Lys | Glu | Gly | Ile | Tyr | Glu | Asn |
| | | | | | | 35 | | 40 | | | | 45 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Ala | Lys | Phe | His | Gly | Lys | Pro | Gly | Leu | Cys | Glu | His | Asp | Gly | His |
| | | | | | 50 | | 55 | | | | 60 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Val | Ala | Cys | Cys | Val | His | Gly | Met | Pro | Thr | Phe | Pro | His | Trp | His |
| | | | | | 65 | | 70 | | | 75 | | | 80 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Leu | Tyr | Val | Leu | Gln | Val | Glu | Asn | Ala | Leu | Leu | Glu | Arg | Gly | Ser |
| | | | | | 85 | | | 90 | | | | 95 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Val | Ala | Val | Pro | Tyr | Trp | Asp | Trp | Thr | Glu | Lys | Ala | Asp | Gly | Ser |
| | | | | | 100 | | | 105 | | | | 110 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Ser | Leu | Ile | Asn | Asp | Ala | Thr | Tyr | Phe | Asn | Ser | Arg | Ser | Gln | Thr |
| | | | | | 115 | | | 120 | | | 125 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Phe | Asp | Pro | Asn | Pro | Phe | Phe | Arg | Gly | His | Ile | Ala | Phe | Glu | Asn | Ala |
| | | | | | 130 | | 135 | | | 140 | | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Thr | Ser | Arg | Asp | Pro | Gln | Pro | Glu | Leu | Trp | Asp | Asn | Lys | Asp | Phe |
| | | | | | 145 | | 150 | | | 155 | | | 160 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Tyr | Glu | Asn | Val | Met | Leu | Ala | Leu | Glu | Gln | Asp | Asn | Phe | Cys | Asp | Phe |
| | | | | | 165 | | | 170 | | | 175 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Ile | Gln | Leu | Glu | Leu | Ile | His | Asn | Ala | Leu | His | Ser | Arg | Leu | Gly |
| | | | | | | 180 | | 185 | | | 190 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Arg | Ala | Lys | Tyr | Ser | Leu | Ser | Ser | Leu | Asp | Tyr | Thr | Ala | Phe | Asp |
| | | | | | 195 | | 200 | | | 205 | | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Val | Phe | Phe | Leu | His | His | Ala | Asn | Val | Asp | Arg | Ile | Trp | Ala | Ile |
| | | | | | 210 | | 215 | | | 220 | | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Trp | Gln | Asp | Leu | Gln | Arg | Tyr | Arg | Lys | Lys | Pro | Tyr | Asn | Glu | Ala | Asp |
| | | | | | 225 | | 230 | | | 235 | | | 240 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Ala | Val | Asn | Glu | Met | Arg | Lys | Pro | Leu | Gln | Pro | Phe | Asn | Asn | Pro |
| | | | | | 245 | | | 250 | | | 255 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Leu | Asn | Ser | Asp | Ser | Met | Thr | Leu | Lys | His | Asn | Leu | Pro | Gln | Asp |
| | | | | | 260 | | | 265 | | | 270 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Phe | Asp | Tyr | Gln | Asn | Arg | Phe | Arg | Tyr | Gln | Tyr | Asp | Asn | Leu | Gln |
| | | | | | 275 | | 280 | | | 285 | | | | | |

70

Phe Asn His Phe Ser Ile Gln Lys Leu Asp Gln Thr Ile Gln Ala Arg
 290 295 300

Lys Gln His Asp Arg Val Phe Ala Gly Phe Ile Leu His Asn Ile Gly
 305 310 315 320

Thr Ser Ala Val Val Asp Ile Tyr Ile Cys Val Glu Gln Gly Gly Glu
 325 330 335

Gln Asn Cys Lys Thr Lys Ala Gly Ser Phe Thr Ile Leu Gly Gly Glu
 340 345 350

Thr Glu Met Pro Phe His Phe Asp Arg Leu Tyr Lys Phe Asp Ile Thr
 355 360 365

Ser Ala Leu His Lys Leu Gly Val Pro Leu Asp Gly His Gly Phe Asp
 370 375 380

Ile Lys Val Asp Val Arg Ala Val Asn Gly Ser His Leu Asp Gln His
 385 390 395 400

Ile Leu Asn Glu Pro Ser Leu Leu Phe Val Pro Gly Glu Arg Lys Asn
 405 410 415

Ile Tyr Tyr

<210> 77

<211> 413

<212> PRT

<213> Megathura crenulata

<400> 77

Asp Gly Leu Ser Gln His Asn Leu Val Arg Lys Glu Val Ser Ser Leu
 1 5 10 15

Thr Thr Leu Glu Lys His Phe Leu Arg Lys Ala Leu Lys Asn Met Gln
 20 25 30

Ala Asp Asp Ser Pro Asp Gly Tyr Gln Ala Ile Ala Ser Phe His Ala
 35 40 45

Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala Ala His Arg His Ala Cys
 50 55 60

Cys Leu His Gly Met Ala Thr Phe Pro Gln Trp His Arg Leu Tyr Thr
 65 70 75 80

Val Gln Phe Glu Asp Ser Leu Lys Arg His Gly Ser Ile Val Gly Leu
 85 90 95

Pro Tyr Trp Asp Trp Leu Lys Pro Gln Ser Ala Leu Pro Asp Leu Val
 100 105 110

Thr Gln Glu Thr Tyr Glu His Leu Phe Ser His Lys Thr Phe Pro Asn
 115 120 125

Pro Phe Leu Lys Ala Asn Ile Glu Phe Glu Gly Glu Gly Val Thr Thr
 130 135 140
 Glu Arg Asp Val Asp Ala Glu His Leu Phe Ala Lys Gly Asn Leu Val
 145 150 155 160
 Tyr Asn Asn Trp Phe Cys Asn Gln Ala Leu Tyr Ala Leu Glu Gln Glu
 165 170 175
 Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu Ile Leu His Asn Gly Ile
 180 185 190
 His Ser Trp Val Gly Gly Ser Lys Thr His Ser Ile Gly His Leu His
 195 200 205
 Tyr Ala Ser Tyr Asp Pro Leu Phe Tyr Ile His His Ser Gln Thr Asp
 210 215 220
 Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Glu His Arg Gly Leu Ser
 225 230 235 240
 Gly Lys Glu Ala His Cys Ala Leu Glu Gln Met Lys Asp Pro Leu Lys
 245 250 255
 Pro Phe Ser Phe Gly Ser Pro Tyr Asn Leu Asn Lys Arg Thr Gln Glu
 260 265 270
 Phe Ser Lys Pro Glu Asp Thr Phe Asp Tyr His Arg Phe Gly Tyr Glu
 275 280 285
 Tyr Asp Ser Leu Glu Phe Val Gly Met Ser Val Ser Ser Leu His Asn
 290 295 300
 Tyr Ile Lys Gln Gln Glu Ala Asp Arg Val Phe Ala Gly Phe Leu
 305 310 315 320
 Leu Lys Gly Phe Gly Gln Ser Ala Ser Val Ser Phe Asp Ile Cys Arg
 325 330 335
 Pro Asp Gln Ser Cys Gln Glu Ala Gly Tyr Phe Ser Val Leu Gly Gly
 340 345 350
 Ser Ser Glu Met Pro Trp Gln Phe Asp Arg Leu Tyr Lys Tyr Asp Ile
 355 360 365
 Thr Lys Thr Leu Lys Asp Met Lys Leu Arg Tyr Asp Asp Thr Phe Thr
 370 375 380
 Ile Lys Val His Ile Lys Asp Ile Ala Gly Ala Glu Leu Asp Ser Asp
 385 390 395 400
 Leu Ile Pro Thr Pro Ser Val Leu Leu Glu Glu Gly Lys
 405 410

<210> 78

<211> 417

<212> PRT

<213> Megathura crenulata

<400> 78

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| His | Gly | Ile | Asn | Val | Arg | His | Val | Gly | Arg | Asn | Arg | Ile | Arg | Met | Glu |
| 1 | | | | 5 | | | | | 10 | | | | 15 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Ser | Glu | Leu | Thr | Glu | Arg | Asp | Leu | Ala | Ser | Leu | Lys | Ser | Ala | Met |
| | | | | | 20 | | | | 25 | | | | 30 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Ser | Leu | Gln | Ala | Asp | Asp | Gly | Val | Asn | Gly | Tyr | Gln | Ala | Ile | Ala |
| | | | | | 35 | | | 40 | | | | 45 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Phe | His | Gly | Leu | Pro | Ala | Ser | Cys | His | Asp | Asp | Glu | Gly | His | Glu |
| | | | | | 50 | | | 55 | | | | 60 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Ala | Cys | Cys | Ile | His | Gly | Met | Pro | Val | Phe | Pro | His | Trp | His | Arg |
| | | | | | 65 | | | 70 | | | 75 | | 80 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Tyr | Thr | Leu | Gln | Met | Asp | Met | Ala | Leu | Leu | Ser | His | Gly | Ser | Ala |
| | | | | | 85 | | | | 90 | | | 95 | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Ala | Ile | Pro | Tyr | Trp | Asp | Trp | Thr | Lys | Pro | Ile | Ser | Lys | Leu | Pro |
| | | | | | 100 | | | 105 | | | 110 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asp | Leu | Phe | Thr | Ser | Pro | Glu | Tyr | Tyr | Asp | Pro | Trp | Arg | Asp | Ala | Val |
| | | | | | 115 | | | 120 | | | 125 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Asn | Asn | Pro | Phe | Ala | Lys | Gly | Tyr | Ile | Lys | Ser | Glu | Asp | Ala | Tyr |
| | | | | | 130 | | | 135 | | | 140 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Thr | Val | Arg | Asp | Pro | Gln | Asp | Ile | Leu | Tyr | His | Leu | Gln | Asp | Glu | Thr |
| | | | | | 145 | | | 150 | | | 155 | | 160 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Thr | Ser | Val | Leu | Leu | Asp | Gln | Thr | Leu | Leu | Ala | Leu | Glu | Gln | Thr |
| | | | | | 165 | | | 170 | | | 175 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asp | Phe | Cys | Asp | Phe | Glu | Val | Gln | Phe | Glu | Val | Val | His | Asn | Ala | Ile |
| | | | | | 180 | | | 185 | | | 190 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| His | Tyr | Leu | Val | Gly | Gly | Arg | Gln | Val | Tyr | Ala | Leu | Ser | Ser | Gln | His |
| | | | | | 195 | | | 200 | | | 205 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Tyr | Ala | Ser | Tyr | Asp | Pro | Ala | Phe | Phe | Ile | His | His | Ser | Phe | Val | Asp |
| | | | | | 210 | | | 215 | | | 220 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Ile | Trp | Ala | Val | Trp | Gln | Ala | Leu | Gln | Lys | Lys | Arg | Lys | Arg | Pro |
| | | | | | 225 | | | 230 | | | 235 | | 240 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Tyr | His | Lys | Ala | Asp | Cys | Ala | Leu | Asn | Met | Met | Thr | Lys | Pro | Met | Arg |
| | | | | | 245 | | | 250 | | | 255 | | | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Phe | Ala | His | Asp | Phe | Asn | His | Asn | Gly | Phe | Thr | Lys | Met | His | Ala |
| | | | | | 260 | | | 265 | | | 270 | | | | |

73

Val Pro Asn Thr Leu Phe Asp Phe Gln Asp Leu Phe Tyr Thr Tyr Asp
 275 280 285

Asn Leu Glu Ile Ala Gly Met Asn Val Asn Gln Leu Glu Ala Glu Ile
 290 295 300

Asn Arg Arg Lys Ser Gln Thr Arg Val Phe Ala Gly Phe Leu Leu His
 305 310 315 320

Gly Ile Gly Arg Ser Ala Asp Val Arg Phe Trp Ile Cys Lys Thr Ala
 325 330 335

Asp Asp Cys His Ala Ser Gly Met Ile Phe Ile Leu Gly Gly Ser Lys
 340 345 350

Glu Met His Trp Ala Tyr Asp Arg Asn Phe Lys Tyr Asp Ile Thr Gln
 355 360 365

Ala Leu Lys Ala Gln Ser Ile His Pro Glu Asp Val Phe Asp Thr Asp
 370 375 380

Ala Pro Phe Phe Ile Lys Val Glu Val His Gly Val Asn Lys Thr Ala
 385 390 395 400

Leu Pro Ser Ser Ala Ile Pro Ala Pro Thr Ile Ile Tyr Ser Ala Gly
 405 410 415

Glu

<210> 79
 <211> 395
 <212> PRT
 <213> Megathura crenulata

<400> 79
 Asp His Ile Ala Gly Ser Gly Val Arg Lys Asp Val Thr Ser Leu Thr
 1 5 10 15

Ala Ser Glu Ile Glu Asn Leu Arg His Ala Leu Gln Ser Val Met Asp
 20 25 30

Asp Asp Gly Pro Asn Gly Phe Gln Ala Ile Ala Ala Tyr His Gly Ser
 35 40 45

Pro Pro Met Cys His Met Xaa Asp Gly Arg Asp Val Ala Cys Cys Thr
 50 55 60

His Gly Met Ala Ser Phe Pro His Trp His Arg Leu Phe Val Lys Gln
 65 70 75 80

Met Glu Asp Ala Leu Ala Ala His Gly Ala His Ile Gly Ile Pro Tyr
 85 90 95

Trp Asp Trp Thr Ser Ala Phe Ser His Leu Pro Ala Leu Val Thr Asp
 100 105 110

His Glu His Asn Pro Phe His His Gly His Ile Ala His Arg Asn Val
 115 120 125
 Asp Thr Ser Arg Ser Pro Arg Asp Met Leu Phe Asn Asp Pro Glu His
 130 135 140
 Gly Ser Glu Ser Phe Phe Tyr Arg Gln Val Leu Leu Ala Leu Glu Gln
 145 150 155 160
 Thr Asp Phe Cys Gln Phe Glu Val Gln Phe Glu Ile Thr His Asn Ala
 165 170 175
 Ile His Ser Trp Thr Gly Gly His Thr Pro Tyr Gly Met Ser Ser Leu
 180 185 190
 Glu Tyr Thr Ala Tyr Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr
 195 200 205
 Asp Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Phe
 210 215 220
 Gln Tyr Asn Ala Ala His Cys Asp Ile Gln Val Leu Lys Gln Pro Leu
 225 230 235 240
 Lys Pro Phe Ser Glu Ser Arg Asn Pro Asn Pro Val Thr Arg Ala Asn
 245 250 255
 Ser Arg Ala Val Asp Ser Phe Asp Tyr Glu Arg Leu Asn Tyr Gln Tyr
 260 265 270
 Asp Thr Leu Thr Phe His Gly His Ser Ile Ser Glu Leu Asp Ala Met
 275 280 285
 Leu Gln Glu Arg Lys Lys Glu Glu Arg Thr Phe Ala Ala Phe Leu Leu
 290 295 300
 His Gly Phe Gly Ala Ser Ala Asp Val Ser Phe Asp Val Cys Thr Pro
 305 310 315 320
 Asp Gly His Cys Ala Phe Ala Gly Thr Phe Ala Val Leu Gly Glu
 325 330 335
 Leu Glu Met Pro Trp Ser Phe Glu Arg Leu Phe Arg Tyr Asp Ile Thr
 340 345 350
 Lys Val Leu Lys Gln Met Asn Leu His Tyr Asp Ser Glu Phe His Phe
 355 360 365
 Glu Leu Lys Ile Val Gly Thr Asp Gly Thr Glu Leu Pro Ser Asp Arg
 370 375 380
 Ile Lys Ser Pro Thr Ile Glu His His Gly Gly
 385 390 395

<210> 80
 <211> 1266
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 80
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 aaggacgtga gtcacctcac ggatgacgag gtgcagtc tccacggcgc cctccatgac 120
 gtcactgcac ctacagggcc tctgagtttca gaagacataa catcttacca tgccgcacca 180
 gctgtgtgt actacaaggc acggaagatc gcctgctgtc tccacggat gcccagttc 240
 cccttctggc acaggcata tgcgtccaa gcccagcggg cactgttgc caaacggaa 300
 actgtcgaa tgccttactg ggactggacg caaacgtga ctcaacttacc atctttgtc 360
 actgaaccca tctacattga cagtaaagggt ggaaagacgtc aaaccaacta ctggtaccgc 420
 ggcgagatacg ctttcatcaa taagaagact ggcgcgactg tagatgatcg cctattcgag 480
 aagggtggagc ctggtcacta cacacatctt atggagactg tcctcgacgc tctcgaacag 540
 gacgaattct gtaaatttga aatccagttc gagttggctc ataatgtat ccattactt 600
 gttggcggta aatttgaata ttcaatgtca aacttggaat acacccctta cgacccccatc 660
 ttcttcctcc accactccaa cgttgcaccgc ctcttcgcca tctggcagcg tcttcaggaa 720
 ctgcgaggaa agaatccaa tgcataatggac tgcacatg aactcgctca ccagcaactc 780
 caacccttca acaggcagc caatccagtc cagtcacaa aggaccactc gacacctgt 840
 gacccctttt attacaaaaca acttggatac agctacgaca gcttaaacctt gaatggaaatg 900
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 agcttccgac tcagtggctt tgggggttct gccaacgtt gttctatgc atgtgtccct 1020
 gatgatgatc cacgcgtga tgactactgc gagaaagcag ggcacttctt cattcttggg 1080
 ggtcaaaagcgtt aatggccgtg gagattctac agacccttct tctatgtat aactgaagcg 1140
 gtacatcacc ttggagttcc gctaagtggc cactactatg tgaaaacaga actttcagc 1200
 gtgaatggca cagcacttcc acctgtatctt ctccctcaac caactgttgc ctaccgaccc 1260
 gggaaa 1266

<210> 81
 <211> 1257
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 81
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 atagatcatt tgactcgtga agaggaatac gagctaagga tggctctgga gagattccag 120
 gcccacacat ccgttgcgtt gtcacggct acagtagagt accatggctt tcctgctcgt 180
 tgcacgcac cagatcaca agtcagggtt gcctgttgc tgcacggcat ggcaccccttc 240
 cctcaactggc accggctgtt cgttacccttgc gtggaaagatg ctcttgcacg gctgtggatcg 300
 cctatcggtt ttccttatttgc ggactggaca aaacctatgtt ctcaccccttcc agacttggca 360
 tcaaatgaga cgtacgtaga cccgtatggc catacacatc ataatccattt cttcaatgca 420
 aatatatctt ttgaggaggg acaccatcac acgagcagga tgatagattt gaaactgttt 480
 gccccagtcg cttttgggaa gcatccatc ctgtttgtat gaaatccgtt cgcattttag 540
 caggaagatt tctgcactt tgagatttgc tttagtttgc tccataattt tattcatgac 600
 tggataggcg gttccgaaga ttactccatg gcccacccgttccatttccatc ctttgacccc 660
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 ccgactgaca tttatgacta cgaggaagtgc tgcacttac gctacgtatc tctaaccgtt 900
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 aataaaccgg ggaaccaacc actcaaaaggctt ggatgttgcatttgc tggtgccaa 1080
 gaaatgcctt gggcgttgc cgcgttgcata aaggtgcataa taactgtactt attgaagaca 1140
 ctccctctcg atgtcgatgg agattatggat gtcactttta aaatttcatgtatgaccc 1200
 aacgcttgc atacggaccc gttccacac gtcacgttgc tttctgagcc agctcac 1257

<210> 86
 <211> 1209
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 86
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 aacaccttga ctaaggctga gaccgacaac ctgagggagg cgctgtgggg tgcgtatggca 120
 gaccacggtc ccaatggctt tcaagctatt gctgcttcc atgaaaacc agctttgtt 180
 cccatgcctg atggccacaa ctactcatgt tgcgtatccg gcatggctac cttccacac 240
 tgcgtatggcc tctacaccaa gcatggagg gatgcaatga gggcgatgg gtctcatgtc 300
 ggccctgcct actgggactg gactgctgcc ttcacccacc tgccaacact ggtcaccgac 360
 acggacaaca accccttcca acatggacac attgattatc tcaatgtcag cacaactcga 420
 tctccccgag acatgctgtt caacgacccc gagcatggat cagagtgcgtt cttctacaga 480
 caagtctct tagctcttga acaaactgtat ttctgcaaat tcgaagttca gtttgagata 540
 acccacaatg ccatccattc ctggacagggt ggccacagcc cctacggat gtccactctc 600
 gacttcaactg cctacgatcc tctcttctgg cttcaccact ccaacaccga cagaatctgg 660
 gctgtctggc aagcttgca agaatacaga ggacttccat acaaccatgc caattgtgag 720
 atccaggcaa tggaaaacgccc cctgaggcct ttcagtgcacg atatcaacca caacccagtc 780
 acaaaggcta acgcgaagcc attagatgtt ttcgagttata atcgggttag cttccagtag 840
 gacaacctca tcttccatgg atacagtatt ccggaaacttgc atcgcgtgct tgaagaaaaga 900
 aaggaggagg acagaatatt tgcgtccctc cttctcaatgtt gatcaagcg tagtgctgt 960
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 aagggtatgtt agcagctaca cctgaggcat gactctgact ttacctttag ggtgaagatt 1140
 gtcggcaccg acgaccacga gcttccttca gacagtgtca aagcaccaac tattgaattt 1200
 gaaccgggc 1209

<210> 87
 <211> 1536
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 87
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 atcaggaaag aagtgtactt cctctccctg caagaggcca acgcaattaa ggatgcactg 120
 tacaagctcc agaatgacga cagtaaagggg ggcttgagg ccatagctgg ctatcacggg 180
 tattcttataa tgcgtccaga aagaggtacc gacaagtatc cctgctgtgt ccacggaaatg 240
 cccgtgttcc cccactggca cccgctgcatt accattcaga tggagagagc tctgaaaaac 300
 catggctctc caatggcat tccttactgg gattggacaa agaagatgtc gagtcttcca 360
 tctttctttg gagattccag caacaacaac ctttctaca aatattacat cggggcgctg 420
 cagcacgaaa caaccaggga cattaatcag agactcttta atcaaaccaa gtttggtaa 480
 tttgattacc tatattacct aactctgcaa gtcctggagg aaaactcgta ctgtgacttt 540
 gaagttcagt atgagatctt ccataacgccc gtccacttcc ggcttggagg aactggaaag 600
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 agtttggata gaatctggat ctttggcag aagttgcaaa agataagaat gaagccttac 720
 tacgcattgg attgtgttgc cgacagactt atgaaaagacc ccctgcattt cttcaactac 780
 gaaaccgtt atgaagatgtt attccccgc atcaacttcc tcccaagcat actgtttgac 840
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 ctgtctcaggc cattcatgtt ccaccgttca gcccattgtgg ctacacgacat cttggtaatc 1260
 ccagtaggtt cggggccatgtt ccttccgcctt aaagtcgtttag taaagagcggg caccaaagtc 1320
 gagtttacac caatagattt gtcggtgaac aaagcaatgg tggagctggg cagctataact 1380

79

gctatggcta aatgcacgt tcccctttc tcttaccacg gcttgaact ggacaaagtc 1440
 tacagcgtcg atcacggaga ctactacatt gctgcaggta cccacgcgtt gtgtgagcag 1500
 aacctcaggc tccacatcca cgtggaacac gagtag 1536

<210> 88

<211> 591

<212> DNA

<213> *Haliotis tuberculata*

<400> 88

ggtcttcgt actgggactg gacgcagcat ctgactcaac tcccagatct ggtgtcagac 60
 ccctgtttg tcgaccggaa aggagggaaag gcccatacaca acgcatacgtt tcgtggaaac 120
 atcaagttt agaataagaa gactgcaaga gctgttgcg atcgcctttt cgagaaggaa 180
 ggaccaggag agaataacccg actctttgaa ggaattctcg atgccttga acaggatgaa 240
 ttctgcact tcgagatcca gttgagttt gctcacaacg ctatccacta cctgggttggc 300
 ggcgtcaca cgtactccat gtctcatctc gactacaccc cctacgaccc cctcttcttc 360
 ctccatcaact ccaacccggaa ccgcatacttc gccatctggg aacgtcttca ggtactcaga 420
 gggaaaggacc ccaacacccgc cgactgcgca cacaacctca tccatgagcc catggaaccc 480
 ttccgtcggc atgagccat ggaaccgttc cgtcggact cgaaccctct tgacctcacc 540
 agggaaaact ccaaaccaat tgacagctt gattatgccc accttggcta c 591

<210> 89

<211> 1245

<212> DNA

<213> *Haliotis tuberculata*

<400> 89

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 catctgacac gcgaggaggat gtacgagctg cgccagacata tggagagatt ccaggccgac 120
 acatccgtt atgggtacca ggctacgggt gaggatcacg gcttacactgc tcgatgtcca 180
 ttcccccggagg ccacaatag gttgcctgt tgcatccacg gcatggcgac attccctcat 240
 tggcacagac tggcgtcact ccaggtggaa gatgctctga tcaggcgagg atgcctata 300
 ggggtccccct actgggactg gactcagccat atggcgcata tccaggact tgccagacaac 360
 gccacctata gagatcccat cagcggggac agcagacaca accccttca cgatgtgaa 420
 gttgcctttg aaaatggacg tacagaacgt caccggatata gtagattttt tgaacaaccc 480
 ttatggca aacatacgcg tcttcgcac agtatagtct atgcttttga gcaggaggac 540
 ttctgcgatt ttgaaggatca atttgagatg acccataata atattcacgc ctggatttgt 600
 ggcggcgaga agtattccat gtcttctcta cactacacag cctcgaccc tatcttctac 660
 cttcgtcact ccaacactga ccggctctgg gcaatttggc aagcgttgca gatacgaaga 720
 aacaggcctt acaaggctca ttgtgttgg tctgaggaaac gccagccctct caaaccttcc 780
 gccttcgtt cccactgaa caacaacgaa aaaacctacg aaaactcggt gcccaccaac 840
 gtttacgact acgaaggaggat ccttggctat actttagatg acctcaactt cggggggcatg 900
 gacctgggtc agcttgagga atacatccag aggcagagac agagagacag gaccttgc 960
 ggttttttc tgcacatcat tggatcatca gcaatgttg aaatcattat agaccatggg 1020
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 ggatttgacc gtttgacaa atatgagatt acagatgaaac tgaggcaact taatctccgt 1140
 gctgatgatg ttttcagcat ctctgttaaa gtaactgatg ttgatggcag tgagctgtcc 1200
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<210> 90

<211> 1251

<212> DNA

<213> *Haliotis assimilis*

<400> 90

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 cccgaggaaa ttaattctct gaggcgggca atggcagacc ttcaatcaga caaaacccgc 120

80

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gggtggatcc agcaaattgc tgctttcac ggggaaccca aatggtgccc aagtcccgat 180
gctgagaaga agttctccctg ctgtgtccat ggaatggctg tttccctca ctggcacaga 240
ctccctgaccg tgcaaggcga gaatgccctg agaaagcatg gatgtctcg agctctcccc 300
tactgggact ggactcggcc cctgtctcac ctacctgatt tggtttggg aagttagcaga 360
actacacccg tgccatattc caccgtggaa gcccggaaacc cctgggtacag cggccatatt 420
gatacaggtg gtgttgacac aacaagaagc gtccgtcaag aactgtatga agtcctggg 480
tttggccatt atactggggt cgctaagcaa gtgcttctgg ctttggagca ggatgacttc 540
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catcatttca acactgacag actctgggt atatggcagg ctctacaaaaa gtacaggggc 720
aaaccttaca attccgccaa ctgcgcatt gcttctatga gaaaacccct acaacccttt 780
ggtctgactg atgagatcaa cccggatgat gagacaagac agcatgctgt tcctttcagt 840
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ctacacatcg gagatagatt cttcatcaga tacgaaggct tttatcttca tggtacaatg 1200
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<210> 91

<211> 1242

<212> DNA

<213> *Haliotis tuberculata*

<400> 91

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cagaacgacg gagtctatga gaatattgcc aagtccacg gcaaggctgg gttgtgtat 180
gataacggtc gcaagggttc ctgttgttc catggaatgc ccaccttccc ccagtggcac 240
aggctctatg tcctccagggt ggagaatgct ttgctggaga gaggatctgc cgtctctgtg 300
ccatactggg actggactga aacatttaca gagctccat ctttgattgc tgaggcttacc 360
tatttcaatt cccgtcaaca aacgtttgac cctaatttctt tcttcagagg taaaatcagt 420
tttggaaatg ctgttacaac acgtgtatccc cagcctgagc tgtaacgttac caggtactac 480
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gagatggttc acaatgttct ccatgttggg ctgggtggaa gagctactta ttcttattttc 600
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ttgtggggca tctggcagga gctgcagagg tacaggaaga agccatcaa tgaagcggat 720
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atttcgtatata gacaacgtca agatcgttg tttgcaggat tcctccatca caacattggg 960
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gacagactct acaggtttga catcagtggaa acactggggg acctcggcat acagctggac 1140
agccatgact ttgacctcgtt catcaagattt caaggagttaa atggatccta ctttgatcca 1200
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<210> 92

<211> 1257

<212> DNA

<213> *Haliotis tuberculata*

<400> 92

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cattcacctg acgggttcca agccattgcc tctttccatg ctctgccacc actctgcctt 180

81

| | | | | | | |
|-------------|------------|-------------|-------------|------------|-------------|------|
| tcaccatctg | cagctcaccg | ttatgcttgc | tgtgtccacg | gcatggctac | atttccccag | 240 |
| tggcacagat | tgtacactgt | acagttccag | gatgcactga | ggagacatgg | agctacggta | 300 |
| ggtgtacccgt | attgggattt | gctgcgaccc | cagtctcacc | taccagagct | tgtcaccatg | 360 |
| gagacatacc | atgatatttq | gagtaacaga | gattttccca | atcctttcta | ccaagccaat | 420 |
| attgagttt | aaggagaaaa | cattacaaca | gagagagaag | tcattgcaga | caaactttt | 480 |
| gtcaaagggt | gacacgtttt | tgataaaactg | gttctcaaa | caagccatcc | tagcgttag | 540 |
| caggaaaact | actgtgactt | tgagattcag | tttggaaattc | ttcacaacgg | cgttcacacg | 600 |
| tgggtcgag | gcagtcgtac | ctactctatac | ggacatcttc | attacgcatt | ctacgaccct | 660 |
| cttttctacc | ttcaccat | ccagacagac | cgtatttggg | caatctggca | agaactccag | 720 |
| gaacagagag | ggctctcg | tgatgaggt | cactgtgctc | tcgagcaat | gagagaacca | 780 |
| ttgaagcctt | tcagcttcg | cgctccttat | aactggaatc | agctcacaca | ggatttctcc | 840 |
| cgaccggagg | acacccctga | ctacagggaa | tttggttatg | aatatgacaa | tttagaattc | 900 |
| ctggaaatgt | cagttgctga | actggatcaa | tacattattt | aacatcaaga | aatatgataga | 960 |
| gtattcgctg | ggttcctgtt | gagtggattt | ggaggttccg | catcagttaa | tttccaggtt | 1020 |
| tgttagagctg | attccacatg | tcaggatgt | gggtacttca | ccgttcttgg | tggcagtgt | 1080 |
| gagatggcgt | gggcatttga | caggcttac | aaatatgaca | ttactgaaac | tctggagaaa | 1140 |
| atgcaccc | gatatgatga | tgacttcaca | atctctgtca | gtctgaccgc | caacaacgg | 1200 |
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<210> 93

<211> 1248

<212> DNA

<213> *Haliotis tuberculata*

<400> 93

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<210> 94

<211> 1206

<212> DNA

<213> *Haliotis tuberculata*

<400> 94

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82

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<210> 95

<211> 1548

<212> DNA

<213> Haliotis tuberculata

<400> 95

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<210> 96

<211> 966

<212> DNA

<213> Megathura crenulata

<400> 96

83

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<210> 97

<211> 1242

<212> DNA

<213> Megathura crenulata

<400> 97

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<210> 98

<211> 1236

<212> DNA

<213> Megathura crenulata

<400> 98

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84

ccctattggg actggacaaa acgaatcgaa catttaccc acctgattc agacgcccact 360
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<210> 99

<211> 1257

<212> DNA

<213> Megathura crenulata

<400> 99

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<210> 100

<211> 1254

<212> DNA

<213> Megathura crenulata

<400> 100

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85

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<211> 510

<212> DNA

<213> Megathura crenulata

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<210> 102

<211> 942

<212> DNA

<213> Megathura crenulata

<400> 102

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<213> *Megathura crenulata*

<210> 104
<211> 1257
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<213> *Megathura crenulata*

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<210> 105
 <211> 1239
 <212> DNA
 <213> Megathura crenulata

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